

## STIC Search Report Biotech-Chem Library

#### STIC Database Tracking Number: 154530

TO: Brian Kwon

Location: REM-3A74/3C70

**Art Unit: 1614** 

Friday, May 27, 2005

Case Serial Number: 09/763616

From: Barb O'Bryen

**Location: Biotech-Chem Library** 

Remsen 1a69

Phone: 571-272-2518

BB

barbara.obryen@uspto.gov

Search Notes

## RUSH



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Access DB# 154530

#### SEARCH REQUEST FORM

#### Scientific and Technical Information Center

equester's Full Name: Brian In Unit: 1614 Phone Numb	Kum	Examiner =	2815	Date: 5	12.6/05	
in Unit: 1614 Phone Numb	per 30	Semal	Number: 69	1/26/	5	
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lease provide a detailed statement of the searc	h topic and describe a	as specifically	as possible the su	bieci matter to	be searched.	
nclude the elected species or structures, keyworthity of the invention. Define any terms that renown. Please attach a copy of the cover sheet,	ords, synonyms, acrony may have a special me	iyms, and regis eaning. Give e	my numbers, and	combine with	the concept or	
itle of Invention: The aporter	(o, pas	and	metho	<u>L</u>		
nventors (please provide full names):	· · · · · · · · · · · · · · · · · · ·					
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Human pep Propionib Novel hum

Peptide # Peptide # Protein # Peptide #

Aam15683 Aam28187 Abb20093

Propionib

Adm03419
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Human sec Peptide # Drosophil Peptide # Peptide # Peptide #

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Human bon Human bra Human liv

Aam55724 Abg49752 Aam03662

Peptide #

ALIGNMENTS

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Chelated zinc finger; therapeutic; treatment; prophylaxis; MPV; mammalian papillomavirus; antiviral; cytostatic; cervical cancer; lesion;
          AAM15683
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ABM36142
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Polysulfide and dithionodisulfide agents, useful for the treatment or prophylaxis of diseases caused by māmmalian papillomavirus, are disruptors of a chelated metal cation domain in an MPV gene encoded Sim MM Ting AE, (MOLE-) INST MOLECULAR & CELL BIOLOGY Tan YJ, Beerheide W, 99WO-AU000724 98AU-00005733 99AU-00001645 (HUGH/) HUGHES E J L. WPI; 2000-256917/22. 03-SEP-1999; 04-SEP-1998; 15-JUL-1999; protein.

Claim 9; Page 50; 78pp; English.

The present invention describes an agent used in the treatment or prophylaxis of a disease caused or exacerbated by MPV (mammalian papillomavirus) comprising a compound capable of reducing, inhibiting or otherwise decreasing the activity of a protein encoded by an MPV gene by facilitating disruption of a chelated metal cation domain present in the protein. An agent of the present invention can be used to treat cervical cancer or its HPV associated precursor lesions or other HPV associated cancers and/or warts. The present sequence represents a specifically claimed conserved motif sequence for a chelated zinc domain

Sequence 37 AA;

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human secreted proteins given in AAA26346 to AAA26458 encode the human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins given in AAY91451 to AAY91691. The human secreted proteins can have activities based on the tissues and cells they are expressed in. Examples of the activities are: cytostatic; antidiabetic; immunosuppressive; antidiation; antidiated in antidiated their; antidiated and their.

Corresponding secreted proteins are useful for preventing, treating or analiotrating medical conditions, e.g. by protein or gene therapy. Also pathological conditions can be diagnosed by determining the amount of the proteins in a sample or by determining the presence of mutations in the polymucleotides, based on which tissues they are most highly expressed in, and include developing products for the diagnosis or treatment of cancer, tumoures, neurodegenerative discaders, developmental abnormalities and foetal deficiencies, blood discrders, discases of the immune system, autoimmune diseases, hepatic and renal diseases inflammation, allergies, Alzheimer's and behavioural disorders, cantionvascular disorders, transplant rejection, diabetes, asthma, sepsis, acne, psoriasis, cardiovascular disorders,
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                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated human genes and the secreted polypeptides they encode, useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders.
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                                                                                                                                                                                                                                                                                                                                     Human secreted protein sequence encoded by gene 17 SEQ ID NO:287.
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Lafleur D, Wei Y, Ni J, Florence KA, Young PE, Brewer
Soppet DR, Endress GA, Ebner R, Olsen HS, Mucenski M;
                                        0, Indels
                                                                               Score 69; DB 3; I
Pred. No. 6.8e+02;
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100.0%;
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37; Conservative
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reproductive disorders, gastrointestinal disorders, respiratory disorders and metabolic disorders. The proteins or polynucleotides can also be used as food additives or preservatives. The proteins are also useful for identifying their binding partners. AAAA26337 to AAAA26345 and AAY91450 are sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                          leukaemia; autoimmune disorder; allergy; cancer; cardiovascular disorder; respiratory disorder; pulmonary disorder; connective tissue disorder; skin disorder; CNS disorder; congenital disorder; infectious disorder; gastrointestinal disorder; human; secreted protein.
                                                                                                                                                                                                                                                                                                  anticonvulsant; antilipaemic; CNS; gynaecological; antiarthritic; antiasthmatic; anti-HIV; virucide; endocrine; cytostatic; immunosuppressive; antiallergic; cardiovascular; respiratory; dermatological; antimicrobial; gastrointestinal; gene therapy; neurodegenerative disease; behavioral disorder; inflammatory condition;
                                                                                                                                                                                                                                                                                                                                                     hyperproliferative disorder; Alzheimer's disease; Parkinson's disease;
Huntington's disease; metabolic disorder; Tay-Sach's disease;
Leash-Whan syndrome; reproductive disorder; immunological disorder;
arthritis; asthma; AIDS; endocrine disorder; immune disorder;
Hodgkin's lymphoma; haematopoietic disorder; muscular disorder;
                                                                                                            Gaps
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                                                                                                                              Novel human secreted protein fragment seqid 294.
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33; Mismatches 0;
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Wei Y;
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LAFLEUR D W.
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RUBEN S M.
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                                                                 Sequence 37 AA;
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The invention describes an isolated human nucleic acid molecule (I)

comprising a polynucleotide having a nucleotide sequence at least 954
identical to: a sequence polynucleotide fragment of ESQ ID NO: X or of
the cDNA sequence included in ATCC Deposit No: Z, which is hybridisable

to SEQ ID NO: X; or a sequence encoding a polypeptide fragment, domain or
epitope of ESQ ID NO: Y or a polypeptide sequence encoded by the cDNA
sequence included in ATCC Deposit No: Z, which is hybridisable to SEQ ID

NO: X, having a biological activity. The nucleic acids and polypeptides,
pharmaceutical formulations and kits are useful in diagnosing and
treating neurodegenerative diseases states, behavioral disorders,
inflammatory conditions, hyperproliferative disorders (e.g. Alzheimer's
disease, Parkinson's disease or Huntington's diseases), metabolic
disorders (e.g. Tay-Sach's disease or Leash-Nyhan syndrome), reproductive
disorders and immune disorders (e.g. Hodgkin's lymphoma), haematopoietic
conditions afflicting connective tissue, skin disorders, allergy,
cancer, cardiovascular, respiratory or pulmonary disorders, allergy,
cancer, pulmonary disorders and gastrointestinal
conditions afflicting connective tissue, skin disorders and gastrointestinal
conditions afflicting connective tissue, skin disorders and gastrointestinal
conditions afflicting owners and sequence does not appear in
the printed specification but is available in electronic format from the
conditions afflicts at ftp.seqd
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New human secreted nucleic acid, useful for diagnosing and treating neurodegenerative, inflammatory, hyperproliferative, metabolic, reproductive, cardiovascular, respiratory or immunological disorders or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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10.8%; Pred. No. 6.8e+02;
ive 33; Mismatches 0;
                                                                                                                                                                                                            Disclosure; SEQ ID NO 294; 234pp; English.
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99US-0123180P.
99US-0125784P.
99US-0126264P.
99US-0126785P.
99US-0126785P.
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25-MAR-1999;
29-MAR-1999;
01-APR-1999;
06-APR-1999;
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05-MAR-1999;
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OUS-0128114P OUS-012845P OUS-0130077P OUS-0130891P OUS-0130891P OUS-0132481P OUS-0132481P OUS-0132481P OUS-0132481P OUS-0132481P OUS-0132481P	99US-01134218P. 99US-01134218P. 99US-0134211P. 99US-0134370P. 99US-0134370P. 99US-01345124P. 99US-0135124P. 99US-0135623P. 99US-0135623P. 99US-0136782P. 99US-013722P. 99US-0137502P. 99US-0137502P. 99US-0137502P. 99US-0137502P. 99US-0137502P. 99US-0138540P. 99US-0138540P. 99US-0138540P. 99US-0138540P.	905 01394558 905 01394587 905 01394587 905 01394587 905 01394608 905 01394608 905 01394628 905 01394628 905 01397598 905 01397598 905 01397598 905 0140338 905 0140338	90S-0141287P 90S-01411842 90S-01421842 90S-0142803P 90S-0142803P 90S-0142803P 90S-01438242 90S-014436242 90S-01440865 90S-01440865 90S-014438318
08-APR-1999 16-APR-1999 19-APR-1999 23-APR-1999 23-APR-1999 23-APR-1999 30-APR-1999 06-MAY-1999 06-MAY-1999	14 - MAX - 1999; 14 - MAX - 1999; 16 - MAX - 1999; 19 - MAX - 1999; 20 - MAX - 1999; 21 - MAX - 1999; 22 - MAX - 1999; 24 - MAY - 1999; 25 - MAX - 1999; 26 - MAY - 1999; 27 - MAY - 1999; 27 - MAY - 1999; 27 - MAY - 1999; 28 - MAY - 1999; 29 - MAY - 1999; 20 - UNN - 1999; 20 - UNN - 1999; 21 - UNN - 1999; 22 - MAY - 1999; 23 - MAY - 1999; 24 - UNN - 1999; 25 - MAY - 1999; 26 - UNN - 1999; 27 - UNN - 1999; 28 - MAY - 1999; 29 - UNN - 1999; 20 - UNN - 1999; 20 - UNN - 1999; 21 - UNN - 1999; 22 - MAY - 1999; 23 - MAY - 1999; 24 - UNN - 1999; 25 - MAY - 1999; 26 - UNN - 1999; 27 - UNN - 1999; 28 - UNN - 1999; 29 - UNN - 1999; 20 - UNN - 1999; 20 - UNN - 1999; 20 - UNN - 1999; 21 - UNN - 1999; 21 - UNN - 1999; 22 - UNN - 1999; 23 - UNN - 1999; 24 - UNN - 1999; 25 - UNN - 1999; 26 - UNN - 1999; 27 - UNN - 1999; 28 - UNN - 1999; 29 - UNN - 1999; 20 - UNN - 1999; 21 - UNN - 1999; 21 - UNN - 1999; 22 - UNN - 1999; 23 - UNN - 1999; 24 - UNN - 1999; 25 - UNN - 1999; 26 - UNN - 1999; 27 - UNN - 1999; 28 - UNN - 1999; 28 - UNN - 1999; 28 - UNN - 1999; 28 - UNN - 1999; 29 - UNN - 1999; 20 - UNN - 1	18 - JUN - 1999 18 - JUN - 1999 22 - JUN - 1999 23 - JUN - 1999 23 - JUN - 1999 24 - JUN - 1999 25 - JUN - 1999 26 - JUN - 1999 27 - JUN - 1999	30 - JUN 1999 01 - JUL 1999 01 - JUL 1999 02 - JUL 1999 08 - JUL 1999 13 - JUL 1999 13 - JUL 1999 14 - JUL 1999 15 - JUL 1999 16 - JUL 1999 16 - JUL 1999 16 - JUL 1999 19 - JUL 1999 19 - JUL 1999 19 - JUL 1999
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The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide expression, comprising any of the 27,400 fully defined nucleotide encoding at least 8 amino acids of any of the 6888 amino acid sequences fully defined in the specification. The probe is a single axon probe that thybridises under high stringency conditions to a nucleic acid molecule acreased in human cells or tissues. Also included are a spatially.

Compression (comprising a plurality of single exon nucleic acid defenses titled above, where each of the plurality of probes for measuring human cells or mucleic acid purples for measuring human gene expression, a vector comprising the single exon addressably isolatable or amplifiable from the plurality, a single comprising and or optionally with conservative amino acid substitutions, and addressably isolatable or mapped appried comprising and expression acids of any of the above-mentioned amino acid contiguous amino acids of any of the above-mentioned amino acid substitutions, and econtiguous amino acids of any of the above-mentioned amino acid substitutions, and econtiguous amino acids of any of the above-mentioned amino acid substitutions, and accomplising and/or licensing single exon probes or microarrays to a customer destring to measure gene expression, a method of providing extended antibody that binds specifically to a peptide cited above.

Contiguous anino measure gene expression of a single exon probe or actored methods of selling and/or licensing single exon probes or microarrays.

Contiguous which contains a database having a plurality of records cited above. The probes may be used as tools for surveying crossession analysis. The probes may be used as tools for surveying crossession analysis. The probes may be used as tools for surveying a crossession analysis. The probes may be used as tools for surveying a lecrations in the genomic locus that indicated essent sequence data for contain part of the probes are used in id
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 69; DB 8; Length 47; 10.8%; Pred. No. 9.4e+02; ive 33; Mismatches 0; Indels
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26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-00608408.
03-AUG-2000; 2000US-00632366.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Pred. No. 9.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human genome derived single exon protein #5622.
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99US-0159295P.
99US-0159329P.
99US-0159331P.
99US-0159637P.
99US-0159638P.
99US-0160741P.
99US-0160776P.
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99US-0160776P.
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99US-0161406P.
99US-0161359P.
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99US-0161920P.
99US-0161992P.
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ABO59388
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Sequence 50 AA;
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                                                                                                                                                The present invention relates to human single exon nucleic acid probes (SENP: see AAI10068-AAI28459). The present sequence is a peptide encoded by one such probe. The SENPs are derived from human Hela cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver.
                                                                                                 Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human cervical epithelial cells.
                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            foetal liver; gene expression; single exon nucleic acid probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoded by human foetal liver single exon probe.
                                                                                                                                                                                                                                                                                   Score 69; DB 4; Length 50;
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                                                                                                                                                                                                                                                                                Query Match
100.0%; Score 69; DB
Best Local Similarity 10.8%; Pred. No. 1e+0
Matches 4; Conservative 33; Mismatches
                                                                                                                              Claim 27; SEQ ID NO 18665; 487pp; English.
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                                                           Rank DR
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                                       (MOLE-) MOLECULAR DYNAMICS INC.
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                                                          Chen W,
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2000US-00608408.
2000US-00632366.
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2000US-0236359P.
2000GB-00024263.
21-SEP-2000; 2000US-0234687P.
        27-SEP-2000; 2000US-0236359P
04-OCT-2000; 2000GB-00024263
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                                                           Hanzel DK,
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                                                                               WPI; 2001-488901/53.
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                                                                                                                                                                                                                                                                50 AA;
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30-JUN-2000;
03-AUG-2000;
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27-SEP-2000;
04-OCT-2000;
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                                                                                                                                                                                                                                                                Sequence
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                                                           Penn SG,
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                                                           The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human focal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a peptide encoded by a single exon nucleic acid probe of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide #283 encoded by probe for measuring placental gene expression.
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Claim 27; SEQ ID NO 25419; 639pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4; Length
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CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       %; Score 69; DB 4;
; Pred. No. 1e+03;
33; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
10.8%; P
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
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30-JUN-2000; 2000US-00608408
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nes 4; Conservative
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                                                                                                                                                                                                                                                                                                                  Human peptide #265 encoded by breast cell single exon nucleic acid probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes.
                                                                                                                                                                                                                                                                                                                                                        Human; microarray; single exon probe; gene expression; breast; disease;
                                     Gaps
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                                   Indels
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                                                                    1e+03;
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               Pred. No. 1e+C
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10.8%; Pre-
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21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
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                                                                                                                                                                                                                                                                               (first entry)
                                   4; Conservative
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           Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                               cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypertension, cardiac arrhythmias and congenital heart disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                      Protein #266 encoded by probe for measuring heart cell gene expression.
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11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
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30-JUN-2000; 2000US-00608408
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27-SEP-2000; 2000US-0236359P
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Best Local Similarity
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                                                                                                                                                                                                                                                                               23-JAN-2002
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                                                                                                            RESULT 10
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RESULT 11 AAM65971

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Gaps

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probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of the probes of the invention
                                                                                                                                                                                                                                                                                                                      Single exon nucleic acid probes for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides a number of single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, liver, cirrhosis, hyperlipoproteinaemia, hyperlipidaemia,
hypercholesterolaemia, coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                          Example 4; SEQ ID NO 25696; 650pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Length 50;
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CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC
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                                                                                                                                                                                                                                                 Rank
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2000US-00608408.
2000US-00632366.
2000US-0234687P.
2000US-0234687P.
                                                                                                                                                                                                              (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                       21-SEP-2000; 2000US-0234687P.
27-SEP-2000; 2000US-0236359P.
04-OCT-2000; 2000GB-00024263.
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                                                                  2000US-0180312P.
2000US-0207456P.
2000US-00608408.
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2000US-0234687P.
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                                 30-JAN-2001; 2001WO-US000667
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                                                                                                                                                                                                                                                 Penn SG, Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                  WPI; 2001-483446/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 50 AA;
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03-AUG-2000; 2
21-SEP-2000; 2
                                                                                                                      03-AUG-2000;
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                                                                  04-FEB-2000;
26-MAY-2000;
                                                                                                        30-JUN-2000;
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09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is a protein encoded by one of the probes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 genome-derived single exon nucleic acid probes useful for analyzing expression in human bone marrow.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia; epilepsy; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human brain expressed single exon probe encoded protein SEQ ID NO: 25696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention provides a number of single exon nucleic acid
                                                                                                                                       Human; bone marrow expressed exon; gene expression analysis; probe; microarray; cancer; leukaemia; lymphoma; myeloma.
                                                                                                        Human bone marrow expressed probe encoded protein SEQ ID NO: 26277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
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AAM65971 standard; protein; 50 AA
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2000US-0207456P.
2000US-00608408.
2000US-00632366.
2000US-0234687P.
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04-OCT-2000; 2000GB-00024263.
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                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-488900/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 50 AA;
                                                                                                                                                                                                                                  WO200157276-A2.
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03-AUG-2000;
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                                   AAM65971;
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Human gene

RESULT 12 AAM5359

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Gaps

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Claim 27; SEQ ID NO 10323; 322pp; English.
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Best Local Similarity 10.8*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-114183/15.
                                                                                                                                                                                                                                             Sequence 50 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a single exon nucleic acid probe (SENP) (I) for measuring human gene expression in a sample derived from human adult liver, comprising one of 13109 defined nucleotide sequences given in the specification (or complements/ fragments). The probe hybridises at high stringency to a nucleic acid molecule expressed in the human adult liver. (I) may be used for predicting, measuring and displaying gene expression in samples derived from human adult liver. The genes identified may be involved in genetic liver diseases such as cirrhosis, hyperlipoproteinaemia, hyperlipidaemia and hypercholesterolaemia which is sesociated with coronary heart disease. ABG47348-ABG59930 represent human liver single exon encoded peptides of the invention. Note: The sequence information for this patent does not appear in the printed specification but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                         Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human adult liver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single exon nucleic acid probe used to measuring gene expression in a human breast.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide #265 encoded by probe for measuring human breast gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe; human; breast disease; breast cancer; development disorder; inflammatory disease; proliferative breast disease; non-carcinoma tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          %; Score 69; DB 4;
; Pred. No. 1e+03;
33; Mismatches
                                                                                                     Claim 27; SEQ ID NO 26284; 658pp; English
                                                                                                                                                                                                                                                                                                                                ftp.wipo.int/pub/published_pct_sequences
Rank DR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-) MOLECULAR DYNAMICS INC
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2000US-00608408.
2000US-00632366.
2000US-0234687P.
2000US-0236359P.
Chen W,
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                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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Hanzel DK,
                            WPI; 2001-488898/53
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                               Sequence 50 AA;
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03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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 Penn
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The present invention relates to novel single exon nucleic acid probes (see AAI00010-AAI10067). The present sequence is a peptide encoded by one auch probe. The probes are useful for measuring human gene expression in a human breast sample, where the probe hybridises at high stringency to a predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human breast, particularly those diseases with polygenic actiology. The diseases include: breast cancer, disorders of development, inflammatory diseases of the breast, fibrocystic changes, proliferative inflammatory diseases of the breast, fibrocystic changes, proliferative this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fib. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human peptide encoded by genome-derived single exon probe SEQ ID 25283.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            measure gene expression in human lung samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 69; DB 4; 10.8%; Pred. No. 1e+03; iive 33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chen W, Rank DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABG35618 standard; peptide; 50 AA
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26-MAY-2000; 2000US-0207456P.
30-JUN-2000; 2000US-0060408.
03-AUG-2000; 2000US-00632366.
21-SEP-2000; 2000US-0234687P.
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The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from thuman lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12819 open reading frames derived from the 12614 probes. Also included are a matroarray comprising the novel set of probes of the numan lung; comprising the novel set of probes which phyridises at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung; measuring gene expression in a sample collection of detectably labeled nucleic acids derived from human lung man, and (b) measuring the labeled nucleic acids derived from human lung array comprising (a) detecting specific hybridiation of each of the acong the acid from early acid from early assigning exons to a single exon probe, labeled nucleic acids from early assigning exons to a single exon probe, and (b) measuring the expression of cach of the exons in several tissues and/or cell types using hybridiation to a single exon microarray having a probe with the exon, where a common pattern of the exons should be assigned to a single pene, comprising (a) identifying exons in the tissues and/or cell types using hybridiation to a single exon microarrays having a probe with the exon, where a common pattern of the exons should be assigned to a single gene, a peptide comprising on the exons in a the tissues and/or cell types using hybridiastion to a single exon in the probe with the exon, where a common pattern of the exons should be assigned to a single gene, a peptide comprising on the exons in a derived man, unique exons in a gene, particularly using human lung derived mRNA and for the study of lung diseases such as astimate (CC analysis, and for identifying exons in a gene, particularly using human lung derived mRNA and for the study of lung diseases (IDD), intertifying exons in a gene ware used for ge 

Sequence 50 AA;

Gaps . Query Match 100.0%; Score 69; DB 5; Length 50; Best Local Similarity 10.8%; Pred. No. 1e+03; Matches 4; Conservative 33; Mismatches 0; Indels

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Search completed: May 27, 2005, 00:32:23

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Appl Appl

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90, Appl 211, App 6, Appli 42, Appl 1, Appli

Sequence S

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Run on:

Searched:

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Result Š

Sequence

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GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew; Stroobant, Paul;
APPLICANT: Goodearl, Andrew; Stroobant, Paul;
APPLICANT: Minghetti, Luisa; Waterfield, Michael; Marchioni, Mark;
APPLICANT: Chen, Maio Su; Hiles, Ian
APPLICANT: Chen, Maio Su; Hiles, Ian
TITLE OF INVENTION: Glial Mitogenic Factors, Their
TITLE OF INVENTION: Preparation and Use
NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 69; DB 4; Length 37; 10.8%; Pred. No. 1e+02; tive 33; Mismatches 0; Indels
US-08-476-509B-46
US-10-125-062-4
US-09-252-991A-23832
US-08-852-666-14
US-09-471-276-941
US-08-01-376-12
US-08-01-376-12
US-08-01-376-12
US-09-2770-767-42140
US-09-2770-767-42140
US-09-2770-767-4313
US-09-2770-767-49134
US-09-419-381-96
US-09-60-793-5
US-09-60-793-5
US-09-60-793-5
US-09-60-793-6
US-09-60-793-6
US-09-60-45-211
US-08-419-381-96
US-09-60-45-211
US-08-410-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: 98 Human Secreted Proteins
TITLE OF INVENTION: 98 Human Secreted Proteins
FILE REFREENCE: PSO31P1
CURRENT APPLICATION NUMBER: US/09/489,847
CURRENT PILING DATE: 2000-01-24
EARLIER PELICATION NUMBER: PCT/US99/17130
EARLIER PILING DATE: 1999-07-29
EARLIER PILING DATE: 1998-07-30
EARLIER PILING DATE: 1998-07-30
EARLIER PILING DATE: 1998-06-5
EARLIER PILING DATE: 1998-08-05
EARLIER PILING DATE: 1998-08-05
EARLIER PILING DATE: 1998-08-05
EARLIER PILING DATE: 1998-08-06
SOOFWARE: PALENTIN VAF. 2.0
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                                                                                                                                                                                                                                                                                                     Sequence 294, Application US/09489847
Patent No. 6476195
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 40, Application US/08036555B Patent No. 5530109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Conservative
    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
  RESULT 2
US-08-036-555B-40
                                                                                                                                                                                                                                                                                          US-09-489-847-294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-489-847-294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 294
LENGTH: 37
    TYPE: PRT
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Sequence 40, Appl
Sequence 46, Appl
Sequence 38, Appl
Sequence 39, Appl
Sequence 91, Appl
Sequence 91, Appl
Sequence 94381, A
Sequence 49598, A
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302, App
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65.762 Million cell updates/sec
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Patent No. 5258
Patent No. 5258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 183,
Sequence 302,
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Sequence 183,
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                                                                                  May 27, 2005, 00:27:20 ; Search time 42 Seconds
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/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
            GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                        US-09-489-847-294
US-08-036-555B-40
US-08-249-322A-40
US-08-469-526A-40
US-08-469-526A-40
US-08-735-021-40
US-08-735-021-40
US-08-735-021-40
US-08-735-021-40
US-08-735-021-40
US-08-735-021-40
US-08-735-06846A-40
US-08-735-06846A-40
US-08-735-06846A-40
US-08-705-196-19
US-08-108-18-38
US-08-691-814B-38
US-08-691-814B-39
US-08-691-814B-39
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US-08-691-814B-39
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US-09-402-181B-302
US-09-721-456-302
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                                                                                                                                                                                                                   513545 segs, 74649064 residues
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5258287-15
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Maximum Match 100%
Listing first 45 summaries
                                                           OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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69
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Match Length
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                                                                                                                                                                                Scoring table:
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                                                                                                                                Title:
Perfect score:
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Gaps

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Sequence 40, Application US/08249322A
Patent No. 5716930
GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew, Stroobant, Paul;
APPLICANT: Minghetti, Luisa, Waterfield, Michael; Marchioni, Mark;
APPLICANT: Chen, Maio Su; Hiles, Ian
TITLE OF INVENTION: Glial Mitogenic Factors, Their
TITLE OF INVENTION: Preparation and Use
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 69; DB 1; Length 39;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                 MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage COMPUTER: IBM OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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                                                                                                                                                                                                                                                      PRIOR APPLICATION: 530
PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PAPLICATION NUMBER: 07/965,173
FILING DATE: 24-MAR-1993
APPLICATION NUMBER: 07/940,389
FILING DATE: 33-0CT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/907,138
FILING DATE: 30-UNY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRL-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 03-ARRIL-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/863,703
FILING DATE: 10-ARRIL-1991
ATTONNEY, AGGNUT INFORMATION:
NAME: TSAI, Christine H.
REFRERENCE/DOCKET NUMBER: LUD 5250.4
TELECOMMUNICATION INFORMATION:
                                                                                                                                  SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,569
FILLIG DATE: 06-UN-1995
CLASSIFCATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York
COUNTRY: USA
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OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
US-08-469-569-40
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Sequence 40, Application US/08469569

Sequence 40, Application US/08469569

GENERAL INPORMATION:

APPLICANT: Goodearl, Andrew; Stroobant, Paul;

APPLICANT: Minghetti, Luisa; Materfield, Michael; Marchioni, Mark;

APPLICANT: Chen, Maio Su; Hiles, Ian

TITLE OF INVENTION: Glal Mitogenic Factors, Their

TITLE OF INVENTION: Preparation and Use

NUMBER OF SEQUENCES: 184

CORRESPONDENCE ADDRESS:

ADDRESSE: Pelfe & Lynch

STREET: 805 Third Avenue

CITY: New York City

STATE. New York

STATE.
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                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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ilarity 10.8%; Pred. No. 1.1e+02;
Conservative 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/036,555B
CLASSIFICATION NUMBER: US/08/036,555B
FILING DATE: 24-MAR-1993
CLASSIFICATION 435
PRIOR APPLICATION NUMBER: 07/965,173
APPLICATION NUMBER: 07/940,389
FILING DATE: 23-0CT-1992
PRIOR APPLICATION DATA: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA: 07/907,138
FILING DATE: 30-UNN-1992
PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRIL-1992
PRIOR PROFINE OF THE OFFICE OFFICE OF THE OFFICE OF THE OFFICE OFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Tsai, Christine H.
REGISTATION NUMBER: 34,266
REFERRICE, DOCKET NUMBER: LUD 5250.4
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (212) 688-9200
TELEFAX: (212) 888-3884
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
CORRESPEDDED TO THE STATE OF THE STREET: ROT THIRD AVENUE CITY: New York City STATE: New York COUNTRY: USA ZIP: 10022
                                                                                                                                                                                                                                                                                                           COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
The 4; Conserve
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US-08-036-555B-40
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Gaps

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0; Indels

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Query Match 100.0%; Score 69; DB 1; Length 39; Best Local Similarity 10.8%; Pred. No. 1.18+02; Matches 4; Conservative 33; Mismatches 0; Indels
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GENERAL INFORMATION:

APPLICANT: Geodearl, Andrew
APPLICANT: Minghetti, Luisa
APPLICANT: Minghetti, Luisa
APPLICANT: Materifield, Michael
APPLICANT: Marchionni, Mark
APPLICANT: Marchionni, Mark
APPLICANT: Chen, Mario
TITLE OF INVENTION: ENEPARATION AND USE
NUMBER OF SEQUENCES: 187
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Processing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 GGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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APPLICATION NUMBER: 08/036,555
FILING DATE: 24-MRA-1933
APPLICATION NUMBER: 07/965,173
APPLICATION NUMBER: 07/940,389
FILING DATE: 33-OCT-1992
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
APPLICATION NUMBER: 07/907,138
FILING DATE: 03-JUN-1992
APPLICATION NUMBER: 07/907,138
FILING DATE: 03-APRL-1992
APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APRL-1992
APPLICATION NUMBER: 0.K. 91 07566.3
FILING DATE: 10-APR-1991
APPLICATION NUMBER: 0.K. 91 07566.3
FILING DATE: 33-APRL-1991
APPLICATION NUMBER: 39,109
RECISTRATION NUMBER: 39,109
REFERENCATION NUMBER: 04585/00200A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM Compatible Pentium
OPERATING SYSTEM: Windows95
SOFTWARE: WordPerfect (Version 7.0)
CURRENT APPLICATION DATE:
APPLICATION WUMBER: US/08/734,591A
FILING DATE: 22-CCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40, Application US/08734591A
Patent No. 5854220
                                      APPLICATION NUMBER: US/08,
FILING DATE: 06 June 1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 39
      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              617-428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-469-526A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boston
Massachusetts
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                                                                FILING DATE: 06
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 22
CLASSIFICATION:
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TELEFAX: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-734-591A-40
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STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 6
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100.0%; Score 69; DB 1; Length 39;
Best Local Similarity 10.8%; Pred. No. 1.1e+02;
Matches 4; Conservative 33; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 CGCCKCCRTTCACRCAGAAGGTCTTCTCTCTCTCTCAGC 39
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
                                                FILING DATE: 26-MAY-1994
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION: 0ATA: 387
FILING DATE: 24-MAR-1993
PRIOR APPLICATION DATA: 389
FILING DATE: 23-OCT-1992
PRIOR APPLICATION DATA: 389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA: 07/907,138
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA: 07/907,138
FILING DATE: 30-JUN-1992
PRIOR APPLICATION DATA: 07/863,703
FILING DATE: 03-APPLIC-1992
PRIOR APPLICATION DATA: 07/863,703
FILING DATE: 03-APPLIC-1991
APPLICATION DATA: 07/865,37
FILING DATE: 03-APPLIC-1991
APPLICATION NUMBER: 07/866,3
FILING DATE: 10-APPLIC-1991
ATTORNEY/AGENT INFORMATION: 0AME: 10-APPLICATION NUMBER: 34,266
REFERENCE/DOCKET NUMBER: 10D 250.4
TELECOMMUNICATION INFORMATION: 0ATA: 0ATA
            омнек: US/08/249,322A
26-MAY-1994
1.
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Patent No. 572849
GENERAL INFORMATION:
APPLICANT: Goodearl, Andrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Stroobant, Paul
APPLICANT: Minghetti, Luisa
APPLICANT: Materfield, Michael
APPLICANT: Marchionni, Mark
APPLICANT: Chen, Maio Su
APPLICANT: Chen, Maio Su
APPLICANT: Chen, Maio Su
TITLE OF INVENTION: ENEPARATION
TITLE OF INVENTION: PREPARATION ON THE PREPARATION OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 176 Federal Street
CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 688-92C
TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-249-322A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 5
US-08-469-526A-40
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: HILES, IAN
APPLICANT: HILES, IAN
TITLE OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR
TITLE OF INVENTION: PREPARATION AND USE
FILE REPERENCE: 04585/00200L
CURRENT APPLICATION NUMBER: US/08/735,021B
CURRENT APPLICATION NUMBER: 08/412,065
BARLIER APPLICATION NUMBER: 08/036,555
EARLIER FILING DATE: 1995-06-06
BARLIER PILING DATE: 1993-03-24
BARLIER PILING DATE: 1993-03-24
BARLIER PILING DATE: 1992-10-23
BARLIER FILING DATE: 1992-06-33
BARLIER FILING DATE: 1992-06-33
BARLIER FILING DATE: 1992-06-33
BARLIER FILING DATE: 1992-06-30
BARLIER PILING DATE: 1992-04-03
NUMBER OF SEQ ID NOS: 192
SOFTWARE: FASTERQ FOR WINGOWS VETSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 69; DB 3;
10.8%; Pred. No. 1.1e+02;
tive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     %; Score 69; DB 2;
; Pred. No. 1.1e+02;
33; Mismatches 0
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/984,085
FILING DATE: 0.1-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/951,747
FILING DATE: 2.5-SEP-1992
FILING DATE: 2.5-SEP-1992
FILING DATE: 0.7-SEP-1992
APPLICATION NUMBER: 07/927,337
FILING DATE: 1.0-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30.162
REFERENCE/DOCKET NUMBER: 04585/017004
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 40, Application US/08735021B Patent No. 6194377 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
10.8%; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: WATERFIELD, MICHAEL
APPLICANT: MARCHIONNI, MARK
APPLICANT: CHEN, MARIO S.
APPLICANT: HILES, IAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 10.8%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: STROOBANT, PAUL APPLICANT: MINGHETTI, LUISA
                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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; ORGANISM: Bos taurus
US-08-735-021-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best_Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear US-08-469-660-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JS-08-735-021-40
                                                                                                                                                                                                                                                                                                                                                                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 40
LENGTH: 39
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0
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US-08-469-660-40
Sequence 40, Application US/08469660
Patent No. 5876973
GENERAL INFORMATION:
APPLICANT: Gwynne, David I.; Marchionni, Mark;
APPLICANT: Moberre N. There of Invention: THEIR PREPARATION AND USE
TITLE OF INVENTION: THEIR PREPARATION AND USE
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 225 Franklin Street
CITY: Boston
STATE: Massachusetts
ZIP: 0211-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 CGCCKCCRITCACRCAGAAGGTCTICTCCCTTCTCAGC 39
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; Pred. No. 1.1e+02;
33; Mismatches 0
                                                                    APPLICATION NUMBER: 08/036,555
FILING DATE: 03-MAR-1993
APLICATION DATA:
APPLICATION NUMBER: 07/965,173
FILING DATE: 23-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/907,138
FILING DATE: 30-UN-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 30-UN-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 03-APR-1992
PRIOR APPLICATION NUMBER: 07/863,703
FILING DATE: 10-APR-1991
ATTORNEY/AGENT INFORMATION:
NAME: 91846F-58-504/, Kitstina
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 04585/00200P
TELEPHONE: (617) 428-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
STORT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/011,396
FILING DATE: 29-JAN-1993
                   08/470,335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 10.8%;
Matches 4; Conservative 33
                                         06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                         FILING DATE: 06-JUN-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0
                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-734-591A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
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APPLICANT: Goodearl, Andrew

APPLICANT: Stroobant, Paul

APPLICANT: Stroobant, Paul

APPLICANT: Winghetti, Luisa

APPLICANT: Waterfield, Michael

APPLICANT: Waterfield, Michael

APPLICANT: Marchionni, Mark

APPLICANT: Hiles, Ian

TITLE OF INVENTION: GLIAL MITOGENIC FACTORS, THEIR

TITLE OF INVENTION: PREPARATION AND USE

NUMBER OF SEQUENCES: 187

CORRESPONDENCE ADDRESS:

ADDRESSEE: Clark . ...

STRDEN
                          3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCCTTCTCAGC 39
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CURKENT AFFLICATION NUMBER: U5/08/734,664A
FILING DATE: 22-OCT-1996
CLASSIFECATION S 56
FRIOR APPLICATION NUMBER: 08/249,322
FILING DATE: 26-MAY-1994
FRIOR APPLICATION NUMBER: 08/036,555
FILING DATE: 24-MAR-1993
FRIOR APPLICATION NUMBER: 07/965,173
FRIOR APPLICATION NUMBER: 07/965,173
FRIOR APPLICATION NUMBER: 07/965,173
FRIOR APPLICATION NUMBER: 07/940,389
FILING DATE: 03-SEP-1992
FRIOR APPLICATION NUMBER: 07/963,703
FRIUNG APPLICATION NUMBER: 07/963,703
FRIUNG APPLICATION NUMBER: 07/863,703
FRIUNG APPLICATION NUMBER: 07/863,703
FRIUNG APPLICATION NUMBER: 07/863,703
FRIUNG APPLICATION NUMBER: UK 91 07566.3
FRIUNG APPLICATION NUMBER: 39.109
REGISTRATION NUMBER: 39.109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04585/002000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: 1BM Compatible Pentium
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (617) 428-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Massachusetts: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 02110
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Boston
                                                                                                                                               -08-734-664A-40
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US-08-734-664A-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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Sequence 40, Application PC/TUS9506846A
GENERAL INFORMATION:
APPLICATE Goodearl, Andrew David; Stroobant, Paul;
APPLICANT: Minghetti, Luisa; Waterfield, Michael; Marchioni, Mark;
APPLICANT: Chen, Maio Su; Hiles, Ian
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  Length 39;
                                      0; Indels
                                                                                                         3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCCTTCTCAGC 39
                                                                               3 CGCCKCCRITCACRCAGAAGGICITCICCTICTCTCAGG 39
Query Match
100.0%; Score 69; DB 3;
Best Local Similarity 10.8%; Pred. No. 1.1e+02;
Matches 4; Conservative 33; Mismatches 0
                                                                                                                                                                                                                                                          APPLICANT: Robert Sklar, Mark Marchionni,
APPLICANT: Bavid I. Gwynne
TITLE OF INVENTION: METHODS FOR ALTERING
TITLE OF INVENTION: MUSCLE CONDITION
NUMBER OF SEQUENCES: 185
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360
MEDIUM TYPE: kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04585/028WO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05083C
FILING DATE: 06-MAY-94
CLASSIFICATION:
                                                                                                                                                                                                                     Sequence 40, Application PC/TUS9405083C; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/209,204
FILING DATE: 08-MAR-94
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-May-93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFRENCE/DOCKET NUMBER: 0458
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                           225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 10.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 225 Frank.
CITY: Boston
STATE: Massachuse
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
PCT-US94-05083C-40
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PCT-US94-05083C-40
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GENERAL INCORNATION:
APPLICANT: Bard, Jonathan A.
APPLICANT: Bard, Jonathan A.
TITLE OF INVENTION: DNA ENCODING GALANN GALR3 RECEPTORS AND
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLp
STREET: 1185 Avenue of The Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 69; DB 3; Length 43; Best Local Similarity 10.8%; Pred. No. 1.3e+02; Matches 4; Conservative 33; Mismatches 0; Indels
NUMBER OF SEQUENCES: 47
CARRESPONDENCE ADDRESS:
ADDRESSER: Howson and Howson
STREET: Spring House Corporate Cutr., PO Box 457
                                                                                                                                COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
COMPUTER: PEDEPTY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/230,196
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 CSRCTNILREPVCLGGCEHIFCSNCVSDCIGTGCPVC 40
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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 02-AUG-1996
FILING DATE: 02-AUG-1996
FILING DATE: 12-FEB-1997
APPLICATION NUMBER: 18 60/038,109
FILING DATE: 19-FEB-1997
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 31,215
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: WST68BUSA
TELECHOME: 215-540-2200
TELECHAN: 215-540-2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,230
FILING DATE: 23-JUL-1997
CLASSIPICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-900-230-46
; Sequence 46, Application US/08900230
; Patent No. 6329197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 43 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: unknown
MOLECULE TYPE: protein
                                                                             Spring House
Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York COUNTRY: U.S.A. ZIP: 11036
                                                                             CITY: Spring Hou
STATE: Pennsylva
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-230-196-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19, Application US/09230196
Patent No. 6307035
GENERAL INFORMATION:
APPLICANT: Rauscher III, Frank J.
APPLICANT: Jensen, David E.
APPLICANT: Jensen, Barkal Associated Protein (BAP-1) and TITLE OF INVENTION: Uses Therefor
  Glial Mitogenic Factors, Their
Preparation and Use
178
                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGG 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Hanson, Norman D.
REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: LUD 5250.5
TELECOMMUNICATION INFORMATION:
TITLE OF INVENTION: Glial Mi-
TITLE OF INVENTION: Preparat.
NUMBER OF SEQUENCES: 178
CORRESPONDENCE ADDRESS:
ADDRESSE: Felfe & Lynch
STREFT: BOS THIR' Avenue
CITY: New York
STATE: New York
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INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
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PCT-US95-06846A-40
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                                                                                                                                           New York
: USA
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                                                                                                                                                           COUNTRY: US
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CITY: Washington
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APPLICANT: Tomasetto, Catherine
APPLICANT: Basset, Paul
APPLICANT: Byrne, Jennifer
TITLE OF INVENTION: Isolated Nucleic Acid Molecules Useful
TITLE OF INVENTION: as Leukemia Markers and in Breast Cancer Prognosis
NUMBER OF SEQUENCES: 124
                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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0
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ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Ave, NW, Suite 600
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: DC
COUNTRY: USA
ZITE: DC
COUNTRY: USA
ZITE: 20005-3934
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/691,814B
FILING DATE: 31-JUL-1996
CLASSIFICATION DATA:
APPLICATION NUMBER: US 60/002,183
FILING DATE: US 09-4UG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 CTACACGCGGGATGCCTGGCTCTTTTGGGGCCCTCGTC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                             0%; Score 69; DB 3;
b; Pred. No. 1.4e+02;
33; Mismatches 0
                                                            52241-C/JPW/ADM
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                NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKGT NUMBER: 52241
TELECOMMUNICATION INFORMATION:
TELEPAN: 212-278-0400
TELEPAX: 212-391-0525
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION
TELEPHONE: 202-371-2600
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TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                   10.8%;
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 10.8%
Matches 4; Conservative
                                                                                                                                                                                LENGTH: 45 amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE:
                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: no
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APPLICANT: Tomasetto, Catherine
APPLICANT: Danasetto, Catherine
APPLICANT: Baset, Paul
APPLICANT: Byrne, Jennifer
TITLE OF INVENTION: Isolated Nucleic Acid Molecules Useful
TITLE OF INVENTION: as Leukemia Markers and in Breast Cancer Prognosis
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kenalon Syrphy
                                                     Gaps
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Query Match 100.0%; Score 69; DB 2; Length 46; Best Local Similarity 10.8%; Pred. No. 1.4e+02; Matches 4; Conservative 33; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Ave, NW, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CONTRET DE CONTRET DE CONTRET DE CONTRET DE CONTRET DE CONTRET READABLE FORM:

MEDLUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE OF DOS/MS-DOS SOFTWALE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/691,814B FILING DATE: J1-UUL-1996 CLASSIFICATION: 435
                                                                                                                                  3 CPKCNEFARTSGDHRPCLKCKCGKTSGGHAEHGCHPC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 CPKCDTFASSGDHFCLKCRCNKTPGGHAEHDGFCKPC 40
                                                                                                  1383.0090001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/002,183
FILING DATE: 09-AUG-1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                             Sequence 39, Application US/08691814B Patent No. 5981218 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Steffe, Eric K.
REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 138
TELECOMMUNICATION: INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: not relevant TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 47 amino acids
TYPE: amino acid
STRANDEDNESS: not relev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 202-371-2543
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
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Sequence 33820, A Sequence 302, App Sequence 302, App

Sequence 183, App

Sequence 4, Appli Sequence 4, Appli Sequence 198399,

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4 US-10-054-611-183
4 US-10-029-386-33820 .
5 US-10-325-810-302
5 US-10-877-146-302
3 US-10-125-062-4
7 US-10-900-857-4
10S-10-424-599-18399
US-10-424-599-18399
US-09-864-761-35391
                                                                                                       US-10-424-599-191604
US-09-864-761-5625
US-10-424-599-15902
S US-10-424-599-15902
7 US-10-926-683-941
5 US-10-424-599-198842
5 US-10-424-599-198892
6 US-10-424-599-19590
6 US-10-424-599-251423
5 US-10-424-599-251423
5 US-10-424-599-251423
6 US-10-424-599-251423
6 US-10-424-599-251423
8 US-10-424-599-251423
9 US-10-424-599-251423
US-10-424-599-251423
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US-10-011-931-15
US-10-011-931-30
US-10-011-931-31
US-10-011-931-33
US-10-011-931-33
US-10-015-931-33
US-10-015-931-33
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US-10-985-299-15
US-10-985-299-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Komatsoulis et al
TITLE OF INVENTION: 98 Human Secreted Proteins
FILE REFERENCE: PZ031P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/351,334
CURRENT FILING DATE: 2003-01-27
PRIOR APPLICATION NUMBER: 60/30,898
PRIOR PILING DATE: 2002-01-25
PRIOR PLING DATE: 2000-01-24
PRIOR PLING DATE: 2000-01-24
PRIOR PLING DATE: 1999-07-29
PRIOR PELING DATE: 1999-07-29
PRIOR PELING DATE: 1999-07-30
PRIOR PELING DATE: 1998-07-30
PRIOR PELING DATE: 1998-07-30
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR PLING DATE: 1998-08-05
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR PLING DATE: 1998-08-15
PRIOR APPLICATION NUMBER: 60/095,496
PRIOR PLING DATE: 1998-08-12
PRIOR PLING DATE: 1998-08-12
PRIOR PLING DATE: 1998-08-16
PRIOR PLING DATE: 1998-08-16
PRIOR PLING DATE: 1998-08-06
PRIOR PLING DATE: 1998-08-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 294, Application US/10351334; Publication No. US20040034196A1; GENERAL INFORMATION:
    US-10-351-334-294
  Sequence 40, Appl
Sequence 40, Appl
Sequence 46, Appl
Sequence 33022, A
Sequence 8, Appli
Sequence 3565, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 294, App
                                                                                         May 27, 2005, 00:35:31 ; Search time 138 Seconds (without alignments) 92.493 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/pubpaa/NSO7_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT_MBW_PUB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-08-736-019-40
US-08-90-230-46
US-08-90-230-46
US-08-900-230-8
US-08-900-230-8
US-08-900-230-8
US-08-64-774-8
5 US-10-347-542-31
US-09-843-676-183
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Maximum Match 100%
Listing first 45 summaries
                                                                 OM protein - protein search, using sw model
                                                                                                                                                                                                   BLOSUM62DX
Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
                                                                                                                                            US-09-763-616-2
69
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Match
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ALIGNMENTS

Length 37;

100.0%; Score 69; DB 15; 10.8%; Pred. No. 5.8e+02;

Query Match Best Local Similarity

NUMBER OF SEQ ID NOS: 376 SOFTWARE: Patentin Ver. 2.0

Q ID NO 294 LENGTH: 37

TYPE: PRT SOFTWARE: SEQ ID NO

ORGANISM: Homo sapiens US-10-351-334-294

Sequence 8, Appli Sequence 31, Appl Sequence 183, App Sequence 183, App Sequence 183, App Sequence 183, App

US-09-438-486-183 US-10-053-758-183 US-10-054-295-183

Sequence 35391, A Sequence 3525, A Sequence 191604, Sequence 24606, Sequence 24606, Sequence 195890, Sequence 195890, Sequence 25143, Sequence 25143, Sequence 2524, A Sequence 2524, A Sequence 32, Appl Sequence 31, Appl Sequence 33, Appl Sequence 114, Appl Sequence 33, Appl Sequence 34, Appl Sequence 36, Appl Sequence 30, Appl

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                             Length 39;
                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Minghetti, Luisa; Waterfield, Michael; M
Chen, Malo Su; Hiles, Lan
TITLE OF INVENTION: Glial Mitogenic Factors, Their
Preparation and Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 69; DB 8;
10.8%; Pred. No. 6.2e+02;
ive 33; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 30-JUN-1992
APPLICATION NUMBER: 07/663,703
FILING DATE: 03-APRIL-1992
APPLICATION NUMBER: U.K. 91 07566.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Tsai, Christine H. REGISTRATION NUMBER: 34,266 REFERENCE/DOCKET NUMBER: LUD 5250.4 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          Sequence 40, Application US/10844218
Publication No. US20050106666A1
GENERAL IRFORMATION:
GRAPH. Goodearl, Andrew; Stroobant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENY APPLICATION DATA:
APPLICATION NUMBER: US/10/844,218
FILING DATE: 12-ANG-2004
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/036,555
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 24-MAR-1993
APPLICATION WUMBER: 07/965,173
APLICATION WUMBER: 23-0CT-1992
APPLICATION NUMBER: 07/940,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 03-SEP-1992
APPLICATION NUMBER: 07/907,138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 10-APRIL-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (212) 838-3884
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INFORMATION FOR SEQ ID NO: 400
SEQUENCE CHARACTERISTICS
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Best Local Similarity 10.8*
Matches 4; Conservative
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                                                                                                             4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM
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                                                      Query Match
Best Local Similarity
Matches 4; Conserv
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US-10-844-218-40
US-08-736-019-40
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      Gaps
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      Indels
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CORRESPONDENCE ADDRESS:
                                                            CDDCQHNTEGQYCQRCKPGFYRDLRRPFSAPDACKPC 37
         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
COMPUTER: IBM Compatible Pentium
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/736,019
FILING DATE: 22-0CT-1996
CLASSIFICATION: 514
PRIOR APPLICATION: 514
PRIOR APPLICATION NUMBER: 08/471,833
FILING DATE: 06-JUN-1995
PRIOR APPLICATION NUMBER: 08/306,555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39,109
ER: 04585/002000
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PRIOR APPLICATION DATE:

PRIOR APPLICATION DATE:

APPLICATION NUMBER:

PRIOR APPLICATION DATE:

APPLICATION NUMBER:

APPLICATION NU
                                                                                                                                                                                                                                                                     Sequence 40, Application US/08736019
Publication No. US20030207799A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Clark & Elbing LLP
176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                         Stroobant, Paul
Minghetti, Luisa
Waterfield, Michael
Marchionni, Mark
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REGISTRATION NUMBER: 39,109
         33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
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(617) 428-7045
                                                                                                                                                                                                                                                                                                                                                                 Goodearl, Andrew
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            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chen, Mario
Hiles, Ian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Massachusetts COUNTRY: U.S.A.
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CITY: Boston
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APPLICANT:
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Publication No. US20010009766A1;
GENERAL INFORMATION:
APPLICANT: Bard, Jonathan A.
TITLE OF INVENTION: UNA ENCODING GALANN GALR3 RECEPTORS AND TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                         Length 47;
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                                         OTHER INFORMATION: MAP TO ACO02053.2

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 6

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 7.4

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 5.7

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 5.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,230
FLING DATE: 23-JUL-1997
CLASSIFICATION: 435
ATTONREY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REGISTRATION INFORMATION:
TELEPECOMMUNICATION INFORMATION:
TELEPEACOMUNICATION INFORMATION:
TELEPEACOMUNICATION 10FORMATION:
TELEPEACOM. 212-278-0400
                                                                                                                                                                                                                                                                                                                                          %; Score 69; DB 14;
; Pred. No. 7.9e+02;
33; Mismatches 0;
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100.0%; Score 69; DB 8;
Best Local Similarity 10.8%; Pred. No. 8.6e+02;
Matches 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Cooper & Dunham LLp
1185 Avenue of The Americas
                                                                                                                                                                                                                         100.0%;
10.8%; P
                                                                                                                                                                                                                                                                  4; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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  ORGANISM: Homo sapiens
                                                                                                                                                                                                                         Query Match
Best Local Similarity
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: U.S.A.
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STATE: New York
COUNTRY: U.S.A.
ZIP: 11036
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                                                                                                                                                                             US-10-029-386-33022
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US-09-864-761-33565
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STREET: 11
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Publication No. US20030194704A1 (
GENREAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Bark, David R.
APPLICANT: Hanzel, DAVID R.
APPLICANT: HOWNER: US/10/029,386
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT PILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 33022
LENGTH: 47
TYPE: PRT
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                                                                            US-08-900-230-46

Sequence 46, Application US/08900230

Sequence 46, Application US/08900230

Publication No. US2001009766A1

GENERAL INFORMATION:

APPLICANT: Bard, Jonathan A.

TITLE OF INVENTION: USES THEREOF

NUMBER OF SEQUENCES: 59

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                %; Score 69; DB 8; Length 45;
; Pred. No. 7.5e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: IBM PC compatible
COMPUTER: BLEM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,230
FILING DATE: 23-UUL-1997
CLASSIFICATION: 435
ATTONIEY/AGENT INPORMATION:
ANALYS ATTONIEST ATTONIEST
ATTONIEY/AGENT INPORMATION:
3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
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                                                                                                                                                                                                                                                                                     ADDRESSEE: Cooper & Dunham LLp
STREET: 1185 Avenue of The Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFRENCE/DOCKET NUMBER: 5224:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-278-0400
TELBFAX: 212-391-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 10.8%;
Matches 4; Conservative 33
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INFORMATION FOR SEQ ID NO: 46
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   New York
: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: si
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US-10-029-386-33022
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Length 53;
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0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |::|:::::::|::|| CTTCATACTTTTCATTCCAATGGGGTGACTTTGCTTC 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 CIRCCGSVHVQSHSRLLTHSRNSIRIGRLIRSECLEC 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Wilson, David
APPLICANT: Wilson, David
APPLICANT: Wo. US20040009530Alk, Steffen
APPLICANT: Wo. US20040009530Alk, Steffen
APPLICANT: Wo. Jiangchun
APPLICANT: Uemura, Robyn
TITLE OF INVENTION: ENGINERED BINDING PROTEINS
FILE REPRENCE: 11001-006-999
CURRENT APPLICATION NUMBER: US/10/347,542
CURRENT PILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 69; DB 15;
10.8%; Pred. No. 9.3e+02;
tive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 100.0%; Score 69; DB 17; Similarity 10.8%; Pred. No. 8.6e+02; 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                     ; Sequence 8, Application US/10864774; Sequence 8, Application No. US2005009646A1; GENERAL INFORMATION: APPLICANT: SULLIVAN, SEAN M.; TILE OF INVENTION: GENE DELIVERY TO TUMORS; FILE REFERENCE: 5853-337-1WO; CURRENT APPLICATION NUMBER: US/10/864,774; CURRENT FILING DATE: 2004-06-09; NUMBER OF SEQ ID NOS: 18; SOFTWARE: PatentIn version 3.2
  33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 183, Application US/09843676
Patent No. US20020164786A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 31, Application US/10347542; Publication No. US20040009530A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Pyrococcus furiosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT; ORGANISM: Rattus norvegicus
US-10-864-774-8
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
US-09-843-676-183
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       4;
                                                                                                                                                                   RESULT 8
US-10-864-774-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 8
LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 31
LENGTH: 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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       Matches
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                                                                         APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR FILE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-X-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: MAP TO APD000091.1

OTHER INFORMATION: EXPRESSED IN BONE MARKOW, SIGNAL = 9

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 12

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 12

OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 7.5

OTHER INFORMATION: EXPRESSED IN BT410, SIGNAL = 7.5

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 9

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 9.5

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 15

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 15

OTHER INFORMATION: SIT_HUMAN HIT: A1891146.1, EVALUE 5.10e+00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR PLILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-02-04
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR PELICATION NUMBER: US 09/632,366
PRIOR PELING DATE: 2000-06-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 09/235,359
PRIOR APPLICATION NUMBER: US 06/236,359
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/174,203 PRIOR FILING DATE: 2001-01-29
         Sequence 33565, Application US/09864761
Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 49117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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LENGTH: 50
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Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: No. US20030032075A1el Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor
                                                                             ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/438,486
FILING DATE: 12-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 CSACSTISGRGAPASWAPLCWAWTISTGPGAPSCCVC 39
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Best Local Similarity 10.8%; Pred. No. 9.5e+02;
Matches 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/724,643
APLING DATE: 01-0CT-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REFERENCE/DOCKET NUMBER: 36,429
Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                 CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US 08/844,419
FILING DATE: 18-APR-1997
                                                                 United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nakamura, Toru
Chapman, Karen B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                     San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
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                                          STATE: Ca
COUNTRY:
ZIP: 9411
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                                                               Andrews, William H.
TITLE OF INVENTION: No. US20020164786Alel Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/843,676
FILING DATE: 26-Apr-2001
CLASSIFICATION: 536
                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Andrews, William H.
TITLE OF INVENTION: No. US20030009019Alel Telomerase
NUMBER OF SUBJUCES: 223
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 CSACSTTSGRGAPASWAPLCWAWTISTGPGAPSCCVC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
-APPLICATION NUMBER: US 08/844,419
                                                                                                                                                                                                                    STATE: California
COUNTRY: United States of America
ZIP: 94111
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
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Publication No. US20030009019A1
GENERAL INFORMATION:
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(415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION TELEPHONE: (415) 576-020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: <Unknown>
  Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 54 amino acids
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Hailey, Calvin
                         Morin, Gregg B.
Harley, Calvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 10.8*
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-438-486-183
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APPLICANT:
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Andrews, William H.
TITLE OF INVENTION: No. US20030059787A1el Telomerase
NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: California
COUNTRY: United States of America
ZIP: 9424A1.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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ADDRESSEE: Townsend and Townsend and Crew STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                        NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REPREBUCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 CSACSTTSGRGAPASWAPLCWAWTISTGPGAPSCCVC 39
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                                                                                                                                              APPLICATION NUMBER: 08/854,050
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/44,419
PILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33; Mismatches
                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/054,295
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APPLICATION NUMBER: US/10/054,611
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/854,050
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Publication No. US2003059787A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (415) 576-0200
                                                                                    FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura, Toru
Chapman, Karen B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 54 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHARACTERISTICS
                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
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Best Local Similarity
Matches 4; Conservat
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US-10-054-611-183
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; Pred. No. 9.5e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: No. US20030044953A1el Telomerase
NUMBER OF SEQUENCES: 225
CORRESPONDENCE ADDRESS:
                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/053,758
FILING DATE: 18-Jan-2002
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/854,050
FILING DATE: 09-MAY-1997
PAPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/44,419
FILING DATE: 18-APR-1997
FILING DATE: 18-APR-1997
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: California
COUNTRY: United States of America
                       COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 183:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMULICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 183, Application US/10054295
Publication No. US20031044953A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO: 183:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
10.8%; E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                              ZIP: 94111
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-053-758-183
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US-10-054-295-183
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Gaps

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Sequence 33820, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David R.
APPLICANT: Hanzel, David R.
APPLICANT: HANSEL, DAVID R.
APPLICANT: HOMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
FILE REFERENCE: AEOMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 69; DB 14; Length 54; Pred. No. 9.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: MAP TO AC002455.1

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.8

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.9

OTHER INFORMATION: SWISSPROT HIT: P34088, EVALUE 1.60e+00
US-10-029-386-33820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: ADD1e, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002930US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |::|::::::::| CSACSTISGEGAPASWAPLCWAWTISTGPGAPSCCVC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | :: | :::::::::| ::: | :: | CPQCWRAQGSHRCMCTVWPRHSVWLWLLVTPICRCC 39
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-007-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 183:
US-10-054-611-183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Search completed: May 27, 2005, 00:48:47 Job time : 149 secs
                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 54 amino acids
                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 183:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
10.8%; P
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Best Local Similarity 10.8%
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Best Local Similarity 10.8*
Matches 4; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 15
US-10-029-386-33820
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LENGTH: 54
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GenCore version 5.1.6

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OM protein - protein search, using sw model

Run on: May 27, 2005, 00:26:25; Search time 39 Seconds

(without alignments)

91.283 Million cell updates/sec
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Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0 Maximum DB seq length: 2000000000 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : PIR 79:\*

1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES Result Query No. Score Match Length DB ID

Description	rubredoxin [valida	rubredoxin - Degul	rubredoxin (valida	rubredoxin [valida	rubredoxin - Helio	probable rubredoxi	rubredoxin - Chlor	٠	rubredoxin - "Buty	rubredoxin (rd-1)	rubredoxin - Therm	rubredoxin - Metha	rubredoxin PAB7224	rubredoxin - Clost	rubredoxin [valida	rubredoxin [valida	rubredoxin [import	rubredoxin - Acine	rubredoxin [import	probable rubA prot	rubredoxin PA5350	. rubredoxin PA5351	testis-specific pr	testis-specific pr	Ġ.	ferredoxin (fdx-8)	rubredoxin XF0379	ferredoxin [import	ferredoxin 2[4Fe-4
ΩI	A33173	JX0241	RUDV	RUDVEG	865620	G71256	A27537	A33182	JU0127	H69359	H72348	B69075	E75136	JU0074	RUCLEP	RUPF	G97241	139520	AE2375	E70593	H82976	A82977	S25774	WTFF	H81133	F69517	B82814	B97333	B42960
DB	-	٦	٦	~	7	7	-	Н	-	~	~	7	~	Н	Н	H	~	~	7	~	7	~	~	М	~	7	7	7	7
Length	52	52	52	52	52	52	53	53	53	53	53	53	53	54	54	54	54	54	54	55	55	55	55	26	56	56	. 57	58	28
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Match	100	100.0	100	100	100	100	100	100	100	100	100.	100.	100.	100.	100.	100.	100.	100.0	100.	100	100	1001	100.	100.0	100.0	100.	100.0	100	100
Score	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69		69
No.	٦	7	m	4	S	9	7	60	D	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	29

ferredoxin 2[4Fe-4	rubredoxin [import	probable rubB prot	ferredoxin (fdx-6)	rubredoxin - Metha	testis-specific pr	hemagglutinin VCA0	ferredoxin - Therm	testis-specific pr	hupl protein - Rhi	hydrogenase-relate	rubredoxin homolog	rubredoxin (rd-2)	testis-specific pr	Mid-ORF2 protein -	anaredoxin [import
FEDV2N	F87558	D70593	B69376	G69073	825772	S18996	A72324	S25775	S27340	853661	E44915	D69418	825773	D36760	AB1988
Н	~	~	Ņ	N	7	N	7	7	N	7	7	7	7	7	7
29	59	9	9	63	63	63	65	68	70	72	72	73	74	74	16
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
69	69	69	69	69	69	69	69	69	69	69	69	69	69	69	69
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

### ALIGNMENTS

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A33173

Cyspecies: Clostridium thermosaccharolyticum
Cyspecies: Clostridium thermosaccharolyticum
Cyspecies: Clostridium thermosaccharolyticum, Clostridium tartarivorum
Cyspecies: Clostridium thermosaccharolyticum, Clostridium tartarivorum
Cyspecies: Clostridium thermosaccharolyticum
Cyspecies: Clostridium thermosaccharolyticum
Cyspecies: J. Gagnon, J.; Sieker, L.C.; Van Dorsselaer, A.; Moulis, J.M.
Biochem. J. 211, 839-841, 1990
A;Title: Rubredoxin from Clostridium thermosaccharolyticum. Amino acid sequence, A;Reference number: A33173; MUD:91058526; PMID:2244884
A;Reference number: A33173; MUD:91058526; PMID:2244884
A;Residues: 1-52 «MEY.
A;Residues: 1-50 «
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6 CTVCGYIYDPEVGDPTQNIPPGTKFEDLPDDWVCPDC 42

8 6

rubredoxin - Desulfovibrio vulgaris (strain Miyazaki)
C;Species: Desulfovibrio vulgaris
C;Species: Desulfovibrio vulgaris
C;Date: 10-Jun-1993 #sequence\_revision 22-Jul-1994 #text\_change 09-Jul-2004
C;Accession: JX0241
R;Shimizu P:, Ogata, M.; Yagi, T.; Wakabayashi, S.; Matsubara, H.
Biochimie 71, 1171-1177, 1989
A;Title: Amino acid sequence and function of rubredoxin from Desulfovibrio vulgaris Miy
A;Reference number: JX0241; MUID:90234754; PMID:2561345
A;Accession: JX0341
A;Molecule type: protein

A; Molecule type: protein
A; Residues: 1-52 <SHI>
A; Cross-references: UNIPROT:P15412
C; Superfamily: rubredoxin; rubredoxin homology
C; Reywords: electron transfer; iron; metalloprotein
F; 3-49/Domain: rubredoxin homology <RUB>
F; 1/Modified site: N-formylmethionine (partial) #status experimental
F; 6,9,39,42/Binding site: iron (Cys) #status predicted

Ouery Match 100.0%; Score 69; DB 1; Length 52; Best Local Similarity 10.8%; Pred. No. 3.1e+02; Matches 4; Conservative 33; Mismatches 0; Indels 0; Gaps

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rubredoxin - Heliobacillus mobilis
C;Species: Heliobacillus mobilis
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
C;Date: 19-Mar-1997 #sequence_revision Biophys 318, 80-88, 1995
Arch. Biochem. Biophys 318, 80-88, 1995
Ayfitle: Isolation, characterization, and primary structure of rubredoxin from the phot Ayfacession: 865620, MuID:95243660, PMID:7726577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:083956; GB:AE001266; GB:AE000520; NID:93323309; PIDN:AAC65:A;Experimental source: strain Nichols
C;Genetics:
                                                                                                                                                                                                                   of the oxidized form at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: protein
A; Molecule type: protein
A; Residues: 1-52 < LEB>
A; Cross-references: UNIPROT: P56263
A; Cross-references: UNIPROT: P56263
C; Comment: Rubredoxin is a nonheme iron protein and substitutes for ferredoxin in some C; Superfamily: rubredoxin; rubredoxin homology
C; Superfamily: rubredoxin; rubredoxin hydroxylation; iron; metalloprotein
F; 3-49/Domain: rubredoxin homology < RUB>
F; 1/Modified site: N-formylmethionine #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the syphilis spirochete.
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R;Fraser, C.M.; Norris, S.J.; Weinstock, G.M.; White, O.; Sutton, G.G.; Dodson, R;Fraser, C.M.; Norris, S.J.; Weinstock, G.M.; White, O.; Sutton, G.G.; Dodson, D.; Howell, J.K.; Chidambaram, M.; Utterback, they, L.; Weidman, J.; Smith, H.O.; Venter, J.C.
Science 281, 375-388, 1998
A;Title: Complete genome sequence of Treponema pallidum, the syphilis spirochete A;Reference number: A71250; WUID:98332770; PMID:966876
A;Accession: G71256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Status: preliminary; nucleic acid sequence not shown; translation not shown
submitted to the Brookhaven Protein Data Bank, March 1988
A;Reference number: A50332; PDB:1RDG
A;Contents: annotation; X-ray crystallography, 1.4 angstroms, residues 1-52
R;Frey, M; Sieker, L; Payan, F; Haser, R; Bruschi, M; Pepe, G; LeGall, J, Mol. Biol. 197, 525-541, 1987
A;Title: Rubredoxin from Desulfovibrio gigas. A molecular model of the oxidi: A;Reference number: A44624; MUID:88155649; PMID:3441010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probable rubredoxin - syphilis spirochete
C;Species: Treponema pallidum subsp. pallidum (syphilis spirochete)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                       A; Contents: annotation; X-ray crystallography, 1.4 angstroms A; Contents: annotation; Trubredoxin homology C; Superfamily: rubredoxin; rubredoxin homology C; Keywords: blocked amino end; electron transfer; iron; metalloprotein F; 3.49/Domain: rubredoxin homology < RUB> F; 1/Modified site: N-formylmethionine #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      F;6,9,39,42/Binding site: iron (Cys) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | | :: | ::::::::| | :: | CLVCGYVYDPAKGDPDHGIAPGTAFEDLPADWVCPLC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |::|::::::::|:::|CTVCGYEYDPAKGDPDSGIKPGTKFEDLPDDWACPVC 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F,6,9,39,42/Binding site: iron (Cys) #status predicted
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10.8%; Pred. No. 3.1e+02;
iive 33; Mismatches 0;
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Pred. No. 3.1e+02;
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33;
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A; Residues: 1-52 <COL>
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c;becies: Desulfovibrio gigas
c;bate: 07-May-1981 #sequence_revision 07-May-1981 #text_change 09-Jul-2004
c;Accession: A00275
R;Bruschi, M.
Biochem: Biophys, Res. Commun. 70, 615-621; 1976
A;Title: The amino acid sequence of rubredoxin from the sulfate reducing bacterium, Desu A;Accession: A00275; MUD:76231572; PMID:938515
A;Moccession: A00275
A;Moccession: L52 < SRUS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession: B33962; A00274 —
R;Brumlik, M.J.; Voordouw, G.
B. Bacteriol. 171, 4996-5004, 1989
A;Title. After the genes for rubredoxin (rub) an A;Reference number: A33962; MUID:89359139; PMID:2549009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bruschi, M.
Jochim. Biophys. Acta 434, 4-17, 1976
Jitle: Non-heme iron proteins. The amino acid sequence of rubredoxin from Desulfovibri
Reference number: A00274; MUID:76232334; PMID:7308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cross-references: UNIPROT: P00269; GB: M28848; NID: 9342028; PIDN: AAA64798.1; PID: 975867
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A; Residues: 1-20, Tr', 22-52 < BRUS>
A; Estidues: 1-20, Tr', 21-52 < BRUS>
A; Estidues: 1.5 Sieker, L.C.; Jensen, L.H.
Submitted to the Brookhaven Protein Data Bank, May 1990
A; Reference number: A50939; PDB: 7RXN
A; Contentes: annotation; X-ray crystallography, 1.5 angstroms, residues 1-52
A; Contentes: annotation; X-ray crystallography, 1.5 angstroms, residues 1-52
B; Adman, E.T.; Seiker, L.C.; Jensen, L.H.
B; Adman, E.T.; Seiker, L.C.; Jensen, L.H.
Adman, E.T.; Seiker, L.S.; Jensen, L.H.
Adman, E.T.; Seiker, L.S.; A resolt
                                                                                                                                                                                                                                                                                                                                                                                           function.
(Species: Desulfovibrio vulgaris
A;Variety: strain Hildenborough
C;Date: 24-Apr-1984 #sequence_revision 30-Jun-1991 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0 angstroms, residues 1-52
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Keywords: electron transfer; iron; metalloprotein
F;3-49/Domain: rubredoxin homology «KUB»
F;6,9,39,42/Binding site: iron (Cys) #status experimental
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                                                                      33; Mismatches
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R;Frey, M.; Sieker, L.C.; Payan, F.
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Best Local Similarity
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R.; GW: T.; MCL

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C;Species: "Butyribacterium methylotrophicum"
C;Species: "Butyribacterium methylotrophicum"
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Date: 31-Mar-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
R;Saeki, K.; Yao, Y.; Wakabayashi, S.; Shen, G.J.; Zeikus, J.G.; Matsubara, H.
J. Biochem. 106, 656-662, 1989
A;Title: Ferredoxin and rubredoxin from Butyribacterium methylotrophicum: complete prim A;Reference number: A91913; MUID:90110065; PMID:2606914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tubredoxin (rd-1) homolog - Archaeoglobus fulgidus
C;Bpecies: Archaeoglobus fulgidus
C;Bpecies: Archaeoglobus fulgidus
C;Bpecies: Archaeoglobus fulgidus
C;Botcies: O5-Dec-1997 #sequence_revision O5-Dec-1997 #text_change O9-Jul-2004
C;Accession: H69359
R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodso.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, B.F Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
Nature 390, 344-370, 1997
A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S. Smith, H.O.; Woses, C.R.; Venter, J.C.
A;Tile: The complete genome sequence of the hyperthermophilic, sulfate-reducing archae A;Reference number: A69250; MUID:98049343; PMID:9389475
A;Accession: H6935
A;Accession: H6935
A;Accession: H6935
A;Accession: A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: DNA
A;Accessions SKLES
A;Access
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C;Comment: Rubredoxin is a nonheme iron protein and substitutes for ferredoxin in some
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C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: H72348
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F;6,9,39,42/Binding site: iron (Cys) #status predicted
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Similarity 10.8%; Score 69; DB 1;
Similarity 10.8%; Pred. No. 3.1e+02;
4; Conservative 33; Mismatches 0
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Best Local Similarity
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Best Local Similarity
Matches 4; Conserv
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Eur. J. Blochem. 163, 161-166, 1987

A,Title: The complete amino acid sequence of rubredoxin from the green phototrophic back A,Note: Chlorobium thiosulphatophilum

A,Note: Chlorobium thiosulphatophilum

A,Accession: A,7537

A,Accession: A,7537

A,Rocioule type: protein

A,Rosidues: 1-53 <a href="Monosorgan">Monosorgan</a>

A,Gross-references: UNIPROT:P09947

C,Superfamily: rubredoxin, rubredoxin homology

C,Superfamily: rubredoxin, rubredoxin metalloprotein

F;3-49/Domain: rubredoxin homology <a href="Monosorgan">RUB</a>

F;6-9,39,42/Binding site: iron (Cys) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rubredoxin - Chlorobium limicola f.sp. thiosulfatophilum
C;Species: Chlorobium limicola f.sp. thiosulfatophilum
C;Date: 05-Jun-1988 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Accession: A27537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C.Species: Clostridium sticklandii
C.Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
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                                                                                                                                                                                  Length 52;
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A;Residues: 1-53 -MEY>
A;Residues: 1-53 -MEY>
A;Cross-references: UNIPROT:P23474
A;Experimental source: ATCC 12662
C;Superfamily: rubredoxin; rubredoxin homology
C;Superfamily: rubredoxin; rubredoxin metalloprotein
F;3-49/Domain: rubredoxin homology akuns
F;1/Modified site: N-formylmethionine #status experimental
                           C;Keywords: iron; metalloprotein
F;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                 CDLCGWGYNPEVGDADGGIPAGTAFENLPDHWECPLC 42
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10.8%; Pred. No. 3.1e+02;
iive 33; Mismatches 0
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10.8%; Pred. No. 3.1e+02;
ive 33; Mismatches 0
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Superfamily: rubredoxin; rubredoxin homology
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R; Meyer, J.; Gagnon, J.; Moulis, J.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rubredoxin - Clostridium sticklandii
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Best Local Similarity 10.8*
Matches 4; Conservative
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A, Reference number: A33182
A, Accession: A33182
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
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Length 53;

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%; Score 69; DB 2; Length 53
; Pred. No. 3.1e+02;
33; Mismatches 0; Indels
A;Gene: rd; PAB7224
C;Superfamily: rubredoxin; rubredoxin homology
C;Rywords: iron; metalloprotein
F;3-49(Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
                                                                                                                                                                                                                                                                                                           |::|::::::::::|| | CKICGYIYDEDEGDPDNGISPGTKFEDLPDDWVCPLC 42
                                                                                                                                                                                                                                                                         100.0%; Score 69;
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Best Local Similarity 10.8*
Matches 4; Conservative
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Best Local S
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C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: B69075
R;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T.;
Olu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, N. Ki, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N.
J. Bacteriol. 179, 7135-7155, 1997
A;Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: funct A;Reference number: A6900; MUID:98037514; PMID:9371463
A;Accession: B69075
A;Accession: B69075
A;Residues: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-53 «MTH>
A;Cross-references: UNIPROT:026259; GB:AE000804; GB:AE000666; NID:g2621196; PIDN:AAB8466
C;Genetics: C;Genetics: Strain Delta H
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A;Cross-references: UNIPROT:Q9V099; GB:AJ248285; GB:AL096836; NID:g5458067; PIDN:CAB498d
A;Experimental source: strain Orsay
C;Genetics:
                                                                                                           :Cross-references: UNIPROT:Q9WZC7; GB:AE001739; GB:AE000512; NID:g4981176; PIDN:AAD3574
Experimental source: strain MSB8
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R;anonymous, Genoscope
submitted to the EMBL Data Library, July 1999
A;Bescription: Pyroccoccus abyssi genome sequence: insights into archaeal chromosome A;Reference number: A75001
A;Reference number: E75136
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C;Species: Methanobacterium thermoautotrophicum
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
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                                                                                                     A.Cross-references: UNIPROT:09WZC7; GB:AE001739; GB:AE0
A.Experimental source: strain MSB8
G.Genetics:
A.Gene: TM0659
C.Superfamily: rubredoxin; rubredoxin homology
C.Keywords: iron; metalloprotein
F;3-49/Pomain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cy8) #status predicted
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C,Superfamily: rubredoxin; rubredoxin homology
C,Keywords: iron; metalloprotein
F;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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10.8%; Pred. No. 3.1e+02;
tive 33; Mismatches 0;
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C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999
C;Accession: E75136
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity Matches 4; Conserv
                            A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-53 <ARN>
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Residues: 1-53 <KAW>
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  A; Accession: H72348
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rubredoxin - Clostridium perfringens
C;Species: Clostridium perfringens
C;Species: Clostridium perfringens
C;Date: 28-Feb-1990 #sequence_revision 22-Jul-1994 #text_change 09-Jul-2004
C;Accession: JU0074
R;Seki, Y.; Seki, S.; Satoh, M.; Ikeda, A.; Ishimoto, M.
A;Title: Rubredoxin from Clostridium perfringens: complete amino acid sequence and part
A;Reference number: JU0074; MUID:90036784; PMID:2553684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPROT:P14072
C;Comment: The protein is reduced with NADH in the presence of a specific NAD(P)H oxido C;Comment: The protein is reduced with homology
C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: electron transfer; iron; metalloprotein
F;3-49/Domain: rubredoxin homology <RUB>
F;6,9,39,42/Binding site: iron (Cys) #status predicted
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A; Residues: 1-13, 'D', 15-21,'D', 23-47,'E', 49-54 < MCC>
A; Residues: 1-13,'D', 15-21,'D', 23-47,'E', 49-54 < MCC>
A; Note: peptides for which chemical overlaps were not determined were positioned on the R; Dauter, Z.; Wilson, K.S.; Sieker, L.C.; Moulis, J.M.; Meyer, J.
submitted to the Brookhaven Protein Data Bank, December 1995
A; Reference number: A6:881; DB1:IRO
A; Contents: annotation; X-ray Drystallography, 1.1 angstroms, residues 1-53
R; Dauter, Z.; Wilson, K.S.; Sieker, L.C.; Moulis, J.M.; Meyer, J.
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Similarity 10.8%; Pred. No. 3.1e+02;
4; Conservative 33; Mismatches 0
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submitted to the Brookhaven Protein Data Bank, December 1995
A;Reference number: A65886; PDB:1IRN
A;Contents: annotation; X-ray crystallography, 1.2 angstroms, residues 1-53
R;Watenpaudh, K.D.; Sieker, L.C.; Jensen, L.H.
J. Mol. Biol. 138, 615-633, 1980
A;Title: Crystallographic refinement of rubredoxin at 1.2 Angstroms resolution.
A;Reference number: A56640; MulD:81005589; PMID:7411618
A;Contents: annotation; X-ray crystallography, 1.2 angstroms
R;Watenpaudh, K.D.; Siecker, L.C.; Herriott, J.R.; Jensen, L.H.
A;Contents: annotation; X-ray crystallography, 1.5 angstrom resolution.
A;Title: Refinement of the model of a protein: rubredoxin at 1.5 angstrom resolution.
A;Title: Refinement of the model of a protein: rubredoxin at 1.5 angstroms
C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: annotation; X-ray crystallography, 1.5 angstroms
C;Superfamily: rubredoxin; rubredoxin homology
C;Keywords: electron transfer; iron; metalloprotein
F;1,Modified site: N-formylmethionine #status experimental
F;6,9,39,42/Binding site: iron (Cys) #status experimental
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6 CTVCGYIYNPEDGDPDNGVNPGTDFKDIPDDWVCPLC 42
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Search completed: May 27, 2005, 00:36:06 Job time : 40 secs THIS PAGE BLANK (USPTO)

clostridium clostridium acinetobact anabaena sp chromobacte

Q9a194 P00268

acinetobact

bacteroides

bordetella bordetella bordetella drosophila

drosophila drosophila mycobacter:

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MEDIATES PROM N.A.

MEDIATRES POSTS 57; PubMed=2510401;

MEDIATRES POSTS 57; PubMed=2510401;

Maloue M.S., Shimomaye E.M.

The completed sequence of lymphocytic choriomeningitis virus reveals
a unique RNA structure and a gene for a zinc finger protein.";

Virology 173:1-10(1989).

Virology 173:1-10(1989).

InterPro; IPR007087; Znf CPH2.

RINE-Pro; IPR007087; Znf CPH2.

RINE-Pro; IPR007087; Znf PP11.

R Pfam; PF03084; Zf-P11; 1.

R ProDom; PD020016; Znf P11; 1.

R PROSITE; PS00028; ZINC_FINGER_C2H2_1; 1.

R NNA-binding; Zinc-finger.
                                                                                                                                                                                                                                                                                                                                                                                                                    Lymphocytic choriomeningitis virus (strain Pasteur).
Viruses; ssRNA negative-strand viruses; Arenaviridae; Arenavirus.
VCBI_TaxID=11625;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Name=TubA; OrderedLocusNames=XCC3268;
Xanthomonas campestris (pv. campestris).
Bacteria; Proteobacteria; Gammaproteobacteria; Xanthomonadaceae; Xanthomonas.
NCBI_TaxID=340;
                                         Q9aqjs
Q8ynk8
Q7plr1
Q7w047
Q7wf74
Q9a4q8
Q01644
Q6qat9
Q6qat9
Q6qat1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5972 MW; 1B2E383153CD8582 CRC64;
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11 CKSCWQKFDSLVRCHDHYLCRHCLNLLLTSSDRCPLC 47
                                                                                                                                                                                                                                                                                                                                           (Rel. 16, Created)
(Rel. 16, Last sequence update)
(Rel. 44, Last annotation update)
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RUBR_ACIAD
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 Local Similarity
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Copyright (c) 1993 - 2005
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InterPro; IPR004039; Rubredox.
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STRATS=ATC 33913

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MEDLINB=22022145; PubMed=12024217; DOI=10.1038/417459a;

A Silva A.C.R., Ferro J.A., Reinach F.C., Farah C.S., Furlan L.R.,

A Quaggio R. B., Monteiro-Vitorello C.B., Van Sluys M.A., Almeida N.F.,

Alves L.M.C., do Amaral A.M., Bertollini M.C., Camargo L.E.A.,

Alves L.M.C., do Amaral A.M., Bertollini M.C., Camargo L.E.A.,

Camarotte G., Cannavan F., Cardozo J., Chambergo F., Ciaphna L.P.,

Camarotte G., Cannavan F., Cardozo J., Chambergo F., Ciaphna L.P.,

Camarotte G., Cannavan F., Cardozo J., Chambergo F., Ciaphna L.P.,

Rarelli R.M.B., Coutinho L.L., Cursino-Santos J.R., El-Dorry H.,

Paria J.B., Ferreira A.J.S., Ferreira R.C.C., Gruber A.,

Ratsuyama A.M., Kishi L.T., Leite R.P., Lemos E.G.M., Lemos M.V.F.,

Locall E.C., Machado M.A., Madeira A.M.B.N., Martine E.C., Marchado M.A., Madeira A.M.B.N., Miyaki C.Y., Moon D.H.,

Martins E.C., Madanis J., Menok C.F.M., Miyaki C.Y., Moon D.H.,

Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,

Spinola L.A.F., Takita M.A., Tandri B., Teixaira B.C., Tezza R.I.D.,

Trindade dos Santos M., Truffi D., Tsai S.M., White F.F.,

Setubal J.C., Kitajima J.P.;

H. "Comparison of the genomes of two Xanthomonas pathogens with differing the per specificities and the setupal decomparison of the setupal decomparison of the setupal decomparison of two Xanthomonas pathogens with differing the per setupal decomparison of the setupal decomparison of two Xanthomonas pathogens with differing the setupal decomparison of two Xanthomonas pathogens with differing the setupal decomparison of two Xanthomonas pathogens with differing the setupal decomparison of the setupal decomparison of two Xanthomonas pathogens with differing the setupal decomparison of the setupal decomparison of two Xanthomonas pathogens with differing the setupal decomparison of the setupal decomparison of two Xanthomonas pathogens with differing the decomparison of two Xanthomonas pathogens with differing the decomparison decomparison decomparis
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STRAIN=306 / ATCC 13902 / XV 101;

MEDLINE=22022145; PubMed=12024217; DOI=10.1038/417459a;

A da Silva A.C.R., Ferro J.A., Reinach F.C., Farah C.S., Furlan L.R.,

A Quaggio R.B., Monteiro-Vitorello C.B., Van Sluys M.A., Almeida N.F.,

A lves L.M.C., do Amaral A.M., Bertolini M.C., Camargo L.E.A.,

A ciaratili R.M.B., Coutinho L.L., Cursinno-Santos J.R., El-borry H.,

Faria J.B., Ferreira A.J.S., Ferreira R.C.C., Ferro M.I.T.,

Katsuyama A.M., Kishi L.T., Leite R.P., Lemos E.G.M., Lemos M.V.F.,

A corali B.C., Machado M.A., Madeira A.M.B.N., Martinez-Rossi N.M.,

Martins E.C., Maidanis J., Menck C.F.M., Miyaki C.Y., Moon D.H.,

Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,

Pereira H.A., Rossi A., Sena J.A.D., Silva C., de Souza R.F.,

A spinola L.A.F., Takita M.A., Tamura R.E., Teixeira E.C., Tezza R.I.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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Bacteria, Proteobacteria, Gammaproteobacteria, Xanthomonadales,
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR004039; Rubredox:
InterPro; IPR001052; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00202; RUBREDOXIN; UNKNOWN_1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
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01-OCT-2002 (TrEMBLrel. 22, Last seq
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SEQUENCE 51 AA; 5679 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   Nature 417:459-463(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI TaxID=92829;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rubredoxin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trindade dos Santos M., Truffi D., Tsai S.M., White F.F., Setubal J.C., Kitajima J.P.; Truffi D., Tsai S.M., White F.F., "Comparison of the genomes of two Xanthomonas pathogens with differing host specificities."; Nature 417:459-463(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=rub2; Synonyms=rbr-2; OrderedLocusNames=CT1101;
Chlorobium tepidum.
Bacteria; Chlorobi; Chlorobia; Chlorobiales; Chlorobiaceae;
Chlorobaculum.
                                                                                                                                                                                                                                                                                                                                                                                                      DB 2; Length 51;
                                                                                                                                                                                                                                                                                                                                                                                           Score 69; DB 2; Level Pred. No. 5.1e+02;
                                                                                                                                                                                                                                                                                                                                                             51 AA; 5706 MW; 6E26E27D93530E8F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CVVCGFLYHEADGIPEEGIAPGTRWQDVPDTWTCPDC 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2003 (Rel. 41, Created)
28-FRB-2003 (Rel. 41, Last sequence update)
28-CCT-2004 (Rel. 45, Last annotation update)
Rubredoxin 2 (Rd 2).
                                                                                                                                                          GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0006118; F:electron transport; IEA.
InterPro; IPR004039; Rubredox:
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin:
Prosing: PF00301; Rubredoxin:
PROSITE; PS00202; RUBREDOXIN; UNKNOWN I.
PROSITE; PS50903; RUBREDOXIN, LIKE; I.
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HSSP; P24297; 1BQ8.
                                                                                                                      EMBL; AE011987; AAM38257.1; -. HSSP; P00268; 1IRO.
                                                                                                                                                                                                                                                                                                                                                                                                          100.08;
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Best Local Similarity
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Matches

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RESULT 5

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"Rubredoxin from Desulfovibrio gigas. A molecular model of the oxidized form at 1.4-A resolution.";
J. Mol. Biol. 197:525-541(1987).
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase.
-!- COFACTOR: Binds 1 iron[III) ion per subunit.
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR; A00275; RUDVEG.
PDB; 1861; NMX; A=1-52.
InterPro; IPR041039; Rubredoxin.
InterPro; IPR041039; Rubredoxin.
                                                                                                                                                                                                                                      Desulfovibrio gigas.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
Desulfovibrionaceae; Desulfovibrio.
                                                                                                                                                                                                                                                                                                                                                                     "The amino acid sequence of rubredoxin from the sulfate reducing bacterium, Desulfovibrio gigas."; Biochem. Biophys. Res. Commun. 70:615-621(1976).
                                                                                                                                                                                                                                                                                                                                                                                                                                       X-RAY CRYSTALLOGRAPHY (1.4 ANGSTROMS).
MEDLLINE=88155649; PubMed=3441010;
Frey M., Sieker L.C., Payan F., Haser R., Bruschi M., Pepe G.,
le Gall J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PEAM; PR00301; RUBREDOXIN.
PRONOTE; PR00163; RUBREDOXIN.
PRODOM; PR001610; RUBREDOXIN.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS00903; RUBREDOXIN; 1.
PROSITE; PS00903; RUBREDOXIN, LIKE; 1.
3D-SETUCTURE; Direct protein sequencing; Electron transport; Formylation; Iron; Metal-binding; Rubredoxin.
MOD_RES 1 1 N-formylmethionine.
DOMAIN 1 52 RUBREDOXIN-like.
                        37
                                         | :: | ::::::::| :: | :: | :: | CTVCGYIYDPEVGDPTQNIPPGTKFEDLPDDWVCPDC 42
                      1 CXXCXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
                                                                                                                                                                   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
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Iron.
Iron.
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                                                                                                                                       STANDARD;
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                                                                                                                                                                                                                       Rubredoxin
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-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- COFACTOR: Binds 1 iron(III) ion per subunit (By similarity).
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR, A33173, A33173.
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                    ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
COMPLETE proteome; Electron transport; Iron; Metal-binding; Bubredoxin. 1 57
                                                                                                                                                                                                                                                        Length 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Firmicutes; Clostridia; Thermoanaerobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 69; DB 1; Length 52;
Pred. No. 5.2e+02;
Mismatches 0; Indels
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Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
4FA4AlB0714E080F CRC64;
                                                                                                                                                                                                     Iron (By similarity).
4239E1BF9C52B203 CRC64,
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Iron (By similarity)
Iron (By similarity)
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Pred. No. 5.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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                                                                                                                                                                                                                                             100.0%; Score o, _
10.8%; Pred. No. 5.2e+
...ive 33; Mismatches
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Similarity 10.8%; Pred. No. 5.2e
4; Conservative 33; Mismatches
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PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
PRINTS; PR00163; RUBREDOXIN.
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InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
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PRINTS; PR00163; RUBREDOXIN.
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Matches 4; Conserv
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05-JUL-2004 (Rel.
Rubredoxin (Rd).
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ID RUBR_CLOTS
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    SOLUTION OF THE STREET 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

PubMed=15077118; DOI=10.1038/nbt959;

PubMed=15077118; DOI=10.1038/nbt959;

PubMed=15077118; DOI=10.1038/nbt959;

Rolonay J.F., Seshadri R., Haveman S.A., Hemme C.L., Paulsen I.T.,

Rolonay J.F., Eisen J.A., Mard N.L., Methe B.A., Brinkac I.M.,

Daugherty S.C., DeBoy R.T., Dodson R.J., Durkin A.S., Madupu R.,

Nelson W.C., Sullivan S.A., Fouts D.E., Haft D.H., Selengut J.,

Peterson J.D., Davidsen T.M., Zafar N., Zhou L., Radune D.,

Dimitrov G., Hance M., Tran K., Khouri H.M., Gill J., Utterback T.R.,

Feldbynum T.V., Wall J. D., Voordouw G., Fraser C.M.;

The genome sequence of the anaerobic, sulfate-reducing bacterium

Desulfovibrio vulgaris Hildenborough.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       X-RAY CRYSTALLOGRAPHY (2 ANGSTROMS).
MEDLINE=7730456; PubMed=881725;
Adman E.T., Sieker L.C., Jensen L.H., Bruschi M., le Gall J.;
"A gtructural model of rubredoxin from Desulfovibrio vulgaris at 2-A
                                                                                                                                                   Gaps
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Biochim. Biophys. Acta 434:4-17(1976).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE-88329747; PubMed=2843442; DOI=10.1016/0378-1119(88)90010-8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales; Desulfovibrionaceae; Desulfovibrio. (CEL_TaxID=882;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=rub; OrderedLocusNames=DVU3184;
Desulfovibrio vulgaris (strain Hildenborough / ATCC 29579 / NCIMB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=76232334; PubMed=7308; DOI=10.1016/0005-2795(76)90030-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rubredoxin (rub) and a putative rubredoxin oxidoreductase (rbo) Desulfovibrio vulgaris Hildenborough."; J. Bacteriol. 171:4996-5004(1989).
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MEDLINE=89359139; PubMed=2549009;
Brumlik M.J., Voordouw G.;
"Analysis of the transcriptional unit encoding the genes for
                                                                                                          Length 52;
                                                                                                                                              Indels
                                                              3E08147A4AC262F7 CRC64;
                                                                                                                                                                                                                 6 CTVCGYEYDDAKGDPDSGIKPGTKFEDLPDDWACPVC 42
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                                                                                                                                                ;
                                                                                                     100.0%; Score 69; DB 1;
10.8%; Pred. No. 5.2e+02;
tive 33; Mismatches 0
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01-FEB-1991 (Rel. 17, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
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MEDLINE=91124457; Pubmed=1992166;
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                                                              5677 MW;
                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                                                                   Query Match
Best Local Similarity
45
46
49
AA;
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P00269;
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                                                                                                 X-YAY CRYSTALLOGRAPHY (0.92 ANGSTROMS).

Dauter Z., Butterworth S., Sieker L.C., Sheldrick G., Wilson K.S.;
Submitted (DEC-1997) to the PDB data bank.

-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                             FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase. COFACTOR: Binds 1 iron(III) ion per subunit. SUBCELLUIAR LOCATION: Cytoplasmic. SIMILARITY: Belongs to the rubredoxin family. SIMILARITY: Contains 1 rubredoxin-like domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Adman E.T., Seiker L.C., Jensen L.H.; "Structure of rubredoxin from Desulfovibrio vulgaris at 1.5-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3D-structure; Complete proteome; Direct protein sequencing; Electron transport; Iron; Metal-binding; Rubredoxin. DOMAIN 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 52;
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. 5.2e+02;
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10.8%; Pred. No. 5.2e
tive 33; Mismatches
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PRINTS; PRODIG1; RUBREDOXIN.
PRODIDM; PRODICE10; RUBREDOXIN.
PROSITE; PSOG202; RUBREDOXIN; 1.
PROSITE; PSSG903; RUBREDOXIN LIKE; 1.
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                                                          J. Mol. Biol. 217:337-352(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rubredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; M81166; AAA23381.1; -
EMBL; AE017320; AAS97654.1; -
PIR; B33962; RUDV.
PDB; 1RB9; X-ray; @=1-52.
PDB; 7KXN; X-ray; @=1-52.
PDB; 8KXN; X-ray; A=1-52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGR, DVU3184; -
InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; M28848; AAA64798.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               M21679; AAA21088.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                    the molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             339
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CONFLICT
STRAND
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Miyazaki F in two crystal forms.";
Acta Crystallogr. D 55:408-413(1999)
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking
acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions
as an electron acceptor and may also stabilize the conformation of

    -!- FUNCTION: Electron acceptor for cytoplasmic lactate dehydrogenase.
    -!- COFACTOR: Binds 1 iron(III) ion per subunit.

                                                                                                                                                                                                                                                                                                                                                                                                      Kitamura M., Koshino Y., Kamikawa Y., Kohno K., Kojima S., Miura K., Sagara T., Akutsu H., Kumagai I., Nakaya T.; "Cloning and expression of the rubredoxin gene from Desulfovibrio vulgaris (Miyazaki F) -- comparison of the primary structure of
                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
STRAIN=Isolate F;
MEDLINE=97236449; PubMed=9116039; DOI=10.1016/S0167-4781(96)00203-5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=90234754; PubMed=2561345; DOI=10.1016/0300-9084(89)90020-5; Shimuzu F., Ogata M., Yagi T., Wakabayashi S., Matsubara H.; "Amino acid sequence and function of rubredoxin from Desulfovibrio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=IAM 12604; PubMed=10089148; DOI=10.1107/S0907444998011810; Misaki S., Morimoto Y., Ogata M., Yagi T., Higuchi Y., Yasuoka N.; "Structure determination of rubredoxin from Desulfovibrio vulgaris
                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria; Deltaproteobacteria; Desulfovibrionales;
Desulfovibrionaceae; Desulfovibrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
3D-structure; Direct protein_sequencing; Electron transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
25-CCT-2004 (Rel. 45, Last annotation update)
Rubredoxin (Rd).
                                                                                                              52 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biochim. Biophys. Acta 1351:239-247(1997).
                                                                                                                                                                                                                                                   Desulfovibrio vulgaris (strain Miyazaki)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             X-RAY CRYSTALLOGRAPHY (1.9 ANGSTROMS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SUBCELLULAR LOCATION: Cytoplasmic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL, D76419; BAA11175.1; -.
PIR; JX0241; JX0241.
PDB; IRDV; X-ray; @=1-52.
INTERPTO; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
Pfam; PP003011; Rubredoxin.
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ProDom; PD001610; Rubredox; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biochimie 71:1171-1177(1989).
                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  desulfoferrodoxin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vulgaris Miyazaki."
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                                                                                                                                                                                                                                                                                                           NCBI_TaxID=883;
                                                                                                              RUBR DESVM
                                                                                                                                                                                                                                   Name=rub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                          RUBR_DESVM
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SEQUENCE, AND MASS SPECTROMETRY.
MEDLINE=95243660; PubMed=7726577; DOI=10.1006/abbi.1995.1207;
Lee W.Y., Brune D.C., Lobrutto R., Blankenship R.E.;
"Isolation, characterization, and primary structure of rubredoxin from the photosynthetic bacterium, Heliobacillus mobilis.";
Arch. Blochem. Biophys. 318:80-88(1995).
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions as an electron acceptor and may also stabilize the conformation of
                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteria; Firmicutes; Clostridia; Clostridiales; Heliobacteriaceae;
Heliobacillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PR00163; RUBREDOXIN.
ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN, 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Direct protein sequencing; Electron transport; Formylation; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the molecule.
-!- COFACTOR: Binds 1 iron(III) ion per subunit (By similarity).
-!- MASS SPECTROMETRY: MW=5671.3; METHOD=MALDI; RANGE=1-52;
                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                              Length 52;
                                                                                                                                                                                                                                                                                           %; Score 69; DB 1; Length 52;
   Pred. No. 5.2e+02;
33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                      6443741A8A8063A2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
PIR; 85520; 855620.
HSSP; P15412; 2RDV.
                                                                                                                                                                                                                                                                                                                                                         |::|:::::::|::|
CTVCGYEYDPAEGDPDNGVKPGTAFEDVPADWVCPIC 42
                                                                                                                                                                                                                                                                                                                                           N-formylmethionine.
Rubredoxin-like.
Metal-binding, Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                               15-UUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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                               Iron.
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Iron.
Iron.
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InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                           100.08;
                                                                                                                                                                                                                                                                      5598 MW;
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Best Local Similarity 10.00
Best Local 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                              42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heliobacillus mobilis.
Formylation; Iron;
                                                                                                                                                                                                                                                                    52 AA;
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01-NOV-1999 (
01-OCT-2003 (
Rubredoxin.
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                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              spirochete.";
Science 281:375-388(1998).
-!- FUNCTION: Rubredoxin is a small nonheme, iron protein lacking
acid-labile sulfide. Its single Fe, chelated to 4 Cys, functions
as an electron acceptor and may also stabilize the conformation of
the molecule (By similarity).
-!- COFACTOR: Binds 1 iron (III) ion per subunit (By similarity).
-!- SIMILARITY: Belongs to the rubredoxin family.
-!- SIMILARITY: Contains 1 rubredoxin-like domain.
                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=Nichols;

MEDLINES-9833770; PubMed=965876; DOI=10.1126/science.281.5375.375;

Fraer C.M., Norris S.J., Weinstock G.M., White O., Sutton G.G.,

Dodson R.J., Gwinn M.L., Hickey E.K., Clayton R.A., Ketchum K.A.,

Sodergren E., Hardham U.M., McLeod M.P., Salzberg S.L., Peterson J.D.,

Khalak H.G., Richardson D.L., Howell J.K., Chidambaram M.,

Utterback T.R., McDonald L.A., Artiach P., Bowman C., Cotton M.D.,

Fujii C., Garland S.A., Hatch B., Horst K., Roberts K.M., Sandusky M.,

Weidman J.F., Smith H.O., Venter J.C.;

"Complete genome sequence of Treponema pallidum, the syphilis
                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                         Treponema pallidum.
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Complete proteome; Electron Transport; Iron; Metal-binding;
Rubredoxin. 1 52 Rubredoxin-like.
                                                                                                  Length 52;
                                                                                                                          0; Indels
                                                             Iron (By similarity):
659241131D3124AD CRC64;
N-formylmethionine.
Rubredoxin-like.
Iron (By similarity).
Iron (By similarity).
                                                    similarity).
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                                                                                                                                                 k; Score 69; DB 1;
    Pred. No. 5.2e+02;
33; Mismatches 0
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                                                                                                                                                                                                                                       52 AA
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(By
                                                    Iron
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InterPro; IPR001052; Rubredoxin.
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                                                                                                                                                                                                                                                               (Rel. 37, Created)
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PRINTS; PR00163; RUBREDOXIN.
                                                                                                  100.08;
                                                                        5619 MW;
                                                                                                              10.8%;
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                                                                                    Query Match
Best Local Similarity 10.0.,
A f Conservative
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                                                                                                                                                                                                                                       STANDARD;
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                                                                                                                                                                                                                                                                                                   Rubredoxin (Rd).
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25-OCT-2004
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  MOD RES
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Gaps
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A Michalak P., Noor M.A.F.;
A Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.

B EMBL, ANS49956; AAS5565.1;
BR GO; GO:0006576; C:extracellular; IEA.

GO; GO:0006952; P:defense response; IEA.

DR GO; GO:0006952; P:response to peet, pathogen or parasite; IEA.

GO; GO:0006913; P:response to peet, pathogen or parasite; IEA.

BR GO; GO:0006913; D:response to peet, pathogen or parasite; IEA.

BR InterPro; IPR001450; 4Fe45 ferredoxin.

BR InterPro; IPR001007; VWF C.

DR PROSITE; PS001208; VWFC I; UNKNOWN 1.

PROSITE; PS001208; VWFC I; UNKNOWN 1.

PROSITE; PS01208; VWFC I; UNKNOWN 1.

PROSITE; PS01208; VWFC I; UNKNOWN 1.
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Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
Clostridium.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name-Mst84Dc; ORFNames=CG17945;
Drosophila mauritiana (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                         DB 1; Length 52;
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to the EMBL/GenBank/DDBJ databases.
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
Iron (By similarity).
A76C9F012A70FIEI CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
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Last annotation update)
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                                                                                                                           Score 69;
                                                                                                                                                                                                                                                                                                                                                                                             Created)
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                                                                                                                           100.08;
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                                                                                 5568 MW;
                                                                                                                                                                                                                                                                                                                                                                                           27,
                                                                                                                                              10.8%;
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                                                                                                                                                                  4; Conservative
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Submitted (APR-1999)
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Best Local Similarity
                                                                                                                     Query Match
Best Local Similarity
    6
39
42
52 AA;
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Schools From 1 122;

STRAIN=ATCC 35405 / DSM 14222;

RubMed=15064399; DOI=10.1073/pnas.0307639101;

Rabendari R., Myers G.S.A., Tettelin H., Elsen J.A., Heidelberg J.F., Dodson R.J.; Davidsen T.M., DeBoy R.T., Fouts D.E., Haft D.H.,

Bodson R.J.; Davidsen T.M., DeBoy R.T., Fouts D.E., Haft D.H.,

Bodson R.J.; Davidsen T.M., Madupu R., Kolonay J.F.,

Call S.A., Davidsen T.M., Madupu R., Kolonay J.F.,

Call S.A., Davidsen T.M., Shetty J., Shartsbeyn A.,

Call S.A., Davidsen T.M., Smajs D., Howell J.K., Pavlsen B.,

Call S.M., McLeod M.P., Smajs D., Howell J.K., Pavlsen B.,

Comparison of the genome of the oral pathogen Treponema denticola or in the other spirochete genomes.";

Proc. Natl. Acad. Sci. U.S.A. 101:5646-5651(2004).

C.I. SIMILARITY: Belongs to the rubredoxin family.

EMBL; AE017249; AASIIS41.1;

RESSP; PO0268; 1B13.
                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treponema denticola.
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Rubredoxin.
OrderadiocusNames-ESC31188;
Geobacter sulfurreducens.
Bacteria; Proteobacteria; Deltaproteobacteria; Desulfuromonadales;
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ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
COMPLete proteome; Electron transport; Iron; Metal-binding; Rubredoxin.
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                        0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
                                                                                      | ... | ....... | .... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... | ... |
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                    33; Mismatches
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InterPro; IPR001052; Rubredoxin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TrEMBLrel. 27,
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                4; Conservative
                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rubredoxin.
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Q747S7;
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Q73NV0
                    Matches
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MEDLINE=21992816; PubMed=11997336; DOI=10.1101/gr.219302;

Bao Q., Tian Y., Li W., Xu Z., Xuan Z., Hu S., Dong W., Yang J.,

Chen Y., Xue Y., Xu Y., Lai X., Huang L., Dong X., Ma Y., Ling L.,

Tan H., Chen R., Wang J., Yu J., Yang H.;

A complete sequence of the T. tengcongensis genome.";

Genome Res. 12:689-700(2002).

-!- SIMILARITY: Belongs to the rubredoxin family.

-!- SIMILARITY: Contains 1 rubredoxin-like domain.

EMBL; AR013163; AAM25316.1; --

HSSP; P00270; IRDG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 69; DB 2; Length 52; 10.8%; Pred. No. 5.2e+02; tive 33; Mismatches 0; Indels
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Bacteria; Firmicutes; Clostridia; Thermoanaerobacteriales;
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                                                                                                                           Guedon E.;
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
--- SIMILARITY: Belongs to the rubredoxin family.
--- SIMILARITY: Contains 1 rubredoxin-like domain.
EMBL; AJ228458; CAB41597.1; ---
HSSP; P00270; IRDG.
                                                                                                                                                                                                                                                                                                                                                        GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA.
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PROSITE; PS50903; RUBREDOXIN LIKE; 1.
Electron transport; Iron; Metal-binding; Rubredoxin.
SEQUENCE 52 AA; 5658 MW; AF0F517D000E2FED CRC64;
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Thermoanaerobacteriaceae; Thermoanaerobacter.
NCBI_TaxID=119072;
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PROSITE; PS50903; RUBREDOXIN LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-UUN-2002 (TrEMBLrel. 21, Created)
01-UUN-2002 (TrEMBLrel. 21, Last seq
01-OCT-2003 (TrEMBLrel. 25, Last anno
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InterPro; IPR001052; Rubredoxin.
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ProDom; PD001610; Rubredox; 1.
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PRINTS; PR00163; RUBREDOXIN.
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[2]
SEQUENCE FROM N.A.
STRAIN=ATCC 35319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Q8R870 Q8R870

RESULT 13 **Q8R870** 

Matches

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0;
                                                 STRAIN=PCA / ATCC 51573;

PubMed=14671304; DOI=10.1126/science.1088727;

PubMed=14671304; DOI=10.1126/science.1088727;

Methe B.A., Nelson K.E., Eisen J.A., Paulsen I.T., Nelson W.C., Heidelberg J.F., Wu D., Wu M., Ward N.L., Beanan M.J., Dodson R.J., Madupu R., Brinkac L.M., Daugherty S.C., DeBoy R.T., Durkin A.S., Gwinn M.L., Kolonay J.F., Sallivan S.A., Haft D.H., Salengut J., Bavidsen T.M., Zafar N., White O., Tran B., Romero C., Forberger H.A., Weidman J.F., Khouri H.M., Felablyum T.V., Utterback T.R., Van Aken S.E., Lovley D.R., Fraser C.M.;

"Genome of Geobacter sulfurreducens: metal reduction in subsurface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ProDom; PD001610; Rubredox; 1.
PROSITE; PS00202; RUBREDOXIN; 1.
PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
Complete proteome; Blectron transport; Iron; Metal-binding; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 69; DB 2; Length 52; 10.8%; Pred. No. 5.2e+02; ive 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5877 MW; 5480469CA72044F6 CRC64;
                                                                                                                                                                                                                                                                         Science 302:1967-1969(2003).
--- SIMILARITY: Belongs to the rubredoxin family.
--- SIMILARITY: Contains 1 rubredoxin-like domain.
BEBL; AE017180; ARN36579.1; --
HSSP; P24297; IUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF00301; Rubredoxin; 1
PRINTS; PR00163; RUBREDOXIN.
Geobacteraceae; Geobacter.
NCBI_TaxID=35554;
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Best Local Similarity 10.8*
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          52 AA;
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Search completed: May 27, 2005, 00:35:22 Job time : 176 secs

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May 27, 2005, 00:30:06 ; Search time 495 Seconds (without alignments) 87.306 Million cell updates/sec
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ptodata/1/paa/US087_COMB.pep:
ptodata/1/paa/US088_COMB.pep:
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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/paa/US091_COMB.pep:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pending Patents AA Main:*

| cgn2_6/ptodata/1/paa/pcTUS COI
| cgn2_6/ptodata/1/paa/USO6_COM
| cgn2_6/ptodata/1/paa/USO8_COM
| cgn2_6/ptodata/1/paa/USO8_COI
| cgn2_6/ptodata/1/paa/USO82_COI
| cgn2_6/ptodata/1/paa/USO82_COI
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| cgn2_6/ptodata/1/paa/USO85_COI
| cgn2_6/ptodata/1/paa/USO86_COI
| cgn2_6/ptodata/1/paa/USO86_COI
| cgn2_6/ptodata/1/paa/USO86_COI
| cgn2_6/ptodata/1/paa/USO89_COI
| cgn2_6/ptodata/1/paa/USO89_COI
| cgn2_6/ptodata/1/paa/USO89_COI
| cgn2_6/ptodata/1/paa/USO99_COI
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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69
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                                                                                                                                                                                                                                                                                                                          Title: .
Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	quence 287	equence 29	33 25 27	22,	40,	0 4	Sequence 40, Appl	6,4	, d	40,	40,	6,4	40,	40,	6,6	, 0	40,	40,	40,	, 4	40,	40,	, ¢	40,	40,	40,	2.4	) 4 5 0	e 40	e 40	96 40	40	e 40	Sequence 40, Appl Sequence 40, Appl	?		
ID	PCT-US99-17130-287	US-10-351-334-294	US-60-273-808-33	PCT-US04-15318-22	US-08-011-396A-40	US-08-209-204A-40	US-08-209-204D-40	US-08-209-204E-40	US-U8-461-097-40.	US-08-466-446-40	US-08-467-459-40	US-08-467-459A-40	US-08-467-459-40 US-08-467-470-40	US-08-467-906-40	US-08-468-731-40	US-U8-468-731A-4U	US-08-469-549-40	US-08-469-549-40	US-08-470-335-40	US-US-4 / U-333B-40 IIS-08-471-148-40	US-08-471-833-40	US-08-471-833-40	US-08-471-855-40	US-08-472-065-40	US-08-472-065F-40	US-08-535-200-40	US-U8-535-200A-40	US-08-734-592-40 US-08-734-665-40	US-08-734-665A-40	US-08-734-665-40	US-08-734-666-40	US-08-736-019-40	US-08-736-019-40	US-08-736-070A-40 US-08-736-070-40		ALIGNMENTS	:/TUS9917130 ences, Inc. in Secreted Proteins PCT/US99/17130 60/094,657 77-30 60/095,486 8-06 60/095,455 8-06 60/095,455 8-06 60/095,455
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Length	37	37	38	9 6	99	9 9	39	39	5 C	9 6	39	о С С	n 61	39	33	א מ	9 6	39	66.0	7 6	36	39	66	n 60	39	99	2,0	ת מ יו	39	39	200	0 0	3.6	8 8 8 8	)		Cion Cion Cion Cion Cion Cion Cion Cion
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TYPE: PRT
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                                                                                                                                                        Query Match 100.0%; Score 69; DB 1; Length 37; Best Local Similarity 10.8%; Pred. No. 3.2e+03; Matches 4; Conservative 33; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: TAN, YEE JOO
APPLICANT: BEERHEIDE, WALTER
APPLICANT: BEERHEIDE, WALTER
APPLICANT: SIM, ANTHONY EUGENE
APPLICANT: SIM, MUI MUI
TITLE OF INVENTION: THERAPEUTIC COMPOUNDS AND METHODS
FILE REPRERNCE: BERN3001/JDB
CURRENT APPLICATION NUMBER: US/09/763,616
CURRENT APPLICATION NUMBER: DOT-05-16
PRIOR APPLICATION NUMBER: PCT/AU99/00724
PRIOR APPLICATION NUMBER: AU PQ1645/99
PRIOR APPLICATION NUMBER: AU PPS733/98
PRIOR FILING DATE: 1999-07-15
PRIOR FILING DATE: 1999-07-15
PRIOR FILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 2
SOSTWARE: PATENTIN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 CDDCQHNTEGQYCQRCKPGFYRDLRRPFSAPDACKPC 37
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100.0%; Pred. No. 3.2e+03;
tive 0; Mismatches 0;
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LOCATION: (35)...(36)
OTHER INFORMATION: Variable amino acid
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LOCATION: (5)...(33)
OTHER INFORMATION: Variable amino acid
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OTHER INFORMATION: Variable amino acid
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/09763616 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: BERNARD, HANS-ULRICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 37; Conservative
NUMBER OF SEQ ID NOS: 364
                                                                                               , ORGANISM: Homo sapiens
PCT-US99-17130-287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: MOD_RES
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US-09-763-616-2
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US-09-763-616-2
                   SOFTWARE: Pa
SEQ ID NO 287
                                                                                TYPE: PRT
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Sequence 294, Application US/10351334 GENERAL INFORMATION:

RESULT 3 US-10-351-334-294 APPLICANT: Komatsoulis et al

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G-PROTEIN COUPLED RECEPTOR
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; Sequence 33, Application US/60278983
; GENERAL INFORMATION:
    APPLICANT: Bristol-Myers Squibb Company
    TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR: TITLE OF INVENTION: HGPRBMY27, EXPRESSED HIGHLY IN THE SPLEEN AND TESTIS
    FILE REFERENCE: D0134.pspl.
    CURRENT APPLICATION NUMBER: US/60/278,983
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APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYNUCLEOTIDE ENCODING A NOVEL HUMAN G-PROTEIN COUPLE
TITLE OF INVENTION: HGPRBMY27, EXPRESSED HIGHLY IN THE SPLEEN AND TESTIS
FILE REFERENCE: D0134
CURRENT FILING DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PATENTING DATE: 3001-03-07
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; Pred. No. 3.2e+03;
33; Mismatches 0;
INVENTION: 98 Human Secreted Proteins
                     CURRENT APPLICATION NUMBER: US/10/351,334
CURRENT APPLICATION NUMBER: US/10/351,334
CURRENT FILING DATE: 2003-01-27
PRIOR APPLICATION NUMBER: 09/489,847
PRIOR PILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: 09/489,847
PRIOR PLING DATE: 2000-01-24
PRIOR PLING DATE: 1999-07-29
PRIOR PLING DATE: 1999-07-29
PRIOR APPLICATION NUMBER: 60/094,657
PRIOR PLING DATE: 1998-06-05
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR APPLICATION NUMBER: 60/095,486
PRIOR PLING DATE: 1998-08-12
PRIOR PLING DATE: 1998-08-15
PRIOR PLING DATE: 1998-08-15
PRIOR PLING DATE: 1998-08-16
PRIOR PLING DATE: 1998-08-16
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILING DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILING DATE: 1998-08-06
PRIOR PLILING DATE: 1998-08-06
PRIOR APPLICATION NUMBER: 60/095,455
PRIOR PLILING DATE: 1998-08-06
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Best Local Similarity 10.8%;
Matches 4; Conservative 33,
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US-60-273-808-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens
US-10-351-334-294
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LENGTH: 38
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100.0%; Score 69; DB 4; Length 39; 10.8%; Pred. No. 3.4e+03; ive 33; Mismatches 0; Indels
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MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 40, Application US/08209204A
GENERAL INFORMATION:
APPLICANT: Robert Sklar, Mark Marchionni,
APPLICANT: David I. Gwynne
TITLE OF INVENTION: METHODS FOR TREATING
NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                       REFERENCE DOCKET NUMBER: 30,162
REFERENCE DOCKET NUMBER: 04585/017004
TELECOMMUNICATION INFORMATION:
TELEBHONE: (617) 549-547-547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04585/028001
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CURRENT APPLICATION DATA.
APPLICATION NUMBER: US/08/209,204A
FILING DATE: 08-MAR-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-MAY-93
PRIOR APPLICATION NUMBER: 08/059,022
                     CLASSIPLCATION: 436
PR.OR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
FILING DATE: 01-DEC-1992
PRIOR APPLICATION NUMBER: 07/951,747
FILING DATE: 25-SEP-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/927,337
FILING DATE: 10-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
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ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
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      29-JAN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4; Conservative
                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Massachusetts
ZIP: 02110-2804
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Best Local Similarity
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US-08-209-204A-40
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Kumar, Manoj
APPLICANT: Christiano, Steven P.
APPLICANT: Christiano, Steven P.
TITLE OF INVENTION: Controlled Release of Active Agents Utilizing Repeat Sequence
TITLE OF INVENTION: Protein Polymers
FILE REPERBUCE: DOC 0077 PB/GC 792-6/DC 5110-PCT
CURRENT APPLICATION NUMBER: PT/US04/15318
CURRENT FILING DATE: 2004-05-14
RIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn version 3.2
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Pred. No. 3.3e+03;
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Sequence 40, Application US/08011396A
GENERAL INFORMATION:
APPLICANT: GWYNNE, David I.; Marchionni, Mark;
APPLICANT: GWYNNE, Robert N.
TITLE OF INVENTION: INHIBITORS OF CELL PROLIFERATION,
TITLE OF INVENTION: THEIR PREPARATION AND USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
CITY: Boston
STATE: Massachusetts
ZIP: 02211-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM
                                                                                                                                                                                                                                                                          2 CAGCAGCGCCCATGTACAACGGGTCGTGCTGCCGC 38
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                                                                                                                                                                                                                                 33; Mismatches
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    2001-03-27
                                                                                                                                                                                      Query Match
Best Local Similarity 10.8%; F
Matches 4; Conservative 33;
CURRENT FILING DATE: 2001-03-2
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
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Matches 4; Conservative
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CURRENT APPLICATION DATA:
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OTHER INFORMATION: rspp
                                                        SEO ID NO 33
LENGTH: 38
TYPE: PRT
ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: unknown
                                                                                                                                                                                                                                                                                                                                                                                               PCT-US04-15318-22
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LENGTH: 39
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Gaps

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APPLICANT: Sklar, Robert
APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
APPLICANT: Gwynne, David
TITLE OF INVENTION: Misorders
TITLE OF INVENTION: Disorders
FILE REFRENCE: 04585/028001
CURRENT APPLICATION NUMBER: US/08/209,204E
CURRENT FILING DATE: 1994-03-08
FRIOR APPLICATION NUMBER: US 08/059,022
PRIOR PELLING DATE: 1993-05-06
NUMBER OF SEQ ID NOS: 420
SEQ ID NO 40
SEQ ID NO 40
LENGTH: 39
       APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: Methods for Treating Muscle Diseases and
TITLE OF INVENTION: Disorders
FILE REFERENCE: 04585/028001
CURRENT APPLICATION NUMBER: US/08/209,204D
CURRENT FILING DATE: 1994-03-18
PRIOR PPLICATION NUMBER: US 08/059,022
PRIOR FILING DATE: 1993-05-06
                                                                                                                                                                                                                                                                                                                               Length 39;
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10.8%; Pred. No. 3.4e+03;
tive 33; Mismatches 0;
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10.8%; Pred. No. 3.4e+03;
iive 33; Mismatches 0;
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APPLICANT: David I Gwynne
TITLE OF INVENTION: METHODS FOR TREATING
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                               NUMBER OF SEQ ID NOS: 417
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-209-204E-40; Sequence 40, Application US/08209204E; GENERAL INFORMATION:
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US-08-461-097-40
; Sequence 40, Application US/08461097
; GENERAL INFORMATION:
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Best Local Similarity 10.00
The Ap Conservative
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Best Local Similarity
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ORGANISM: Bos taurus
US-08-209-204E-40
                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Bos taurus
                                                                                                                                                                                                                                                                                                US-08-209-204D-40
                                                                                                                                                                                                                                         LENGTH:
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                                                                                                                                                                                   METHODS FOR TREATING MUSCLE
DISEASES AND DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OCMENTER: IBM Compatible
OPERATING SYSTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/209,204B
FILING DATE: 08-MAR-1994
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   %; Score 69; DB 6;
; Pred. No. 3.4e+03;
33; Mismatches 0
                                                                                                          %; Score 69; DB 6;
; Pred. No. 3.4e+03;
33; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 06-MAY-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 04585,
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
TELEFAX: 617-428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Clark & Elbing LLP
176 Federal Street
                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sklar, Robert
APPLICANT: Marchinoni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: METHODS FO
TITLE OF INVENTION: DISBASES I
NUMBER OF ENGURNES: 379
CORRESPONDENCE ADDRESS:
                                                                                                          100.0%;
10.8%; F
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Best Local Similarity 10.8%; Matches 4; Conservative 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 39 amino acids
                                                                                                                                                 Conservative
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               TYPE: amino acid
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STRANDEDNESS:
;
TOPOLOGY: linear
US-08-209-204A-40
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                                                                                                          Query Match
Best Local Similarity
Matches 4; Conserv
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STATE: MA
COUNTRY: USA
LENGTH: 39
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CITY: Boston STATE: Massachusetts 71P: 02110-2804 ZIP: 02110-2804 COMPUTER READABLE FORM:

US-08-209-204D-40; Sequence 40, Application US/08209204D; GENERAL INFORMATION:

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GENERAL INFORMATION:
APPLICANT: GWynne, David I.; Marchionni, Mark;
APPLICANT: McBurney, Robert N.
TITLE OF INVENTION: INHIBITORS OF CELL PROLIFERATION,
TITLE OF INVENTION: THEIR PREPARATION AND USE
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Boston
STATE: Massachusetts
ZIP: 0211-2804
COMPUTER RRADABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                               3 CGCCKCCRITCACRCAGAAGGTCTICTCCTICTCAGC 39
                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 04585/028002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 04585/017004
FELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PILING DATE:
CLASSIFICATION: 436
PILING DATE:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 29-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/984,085
PILING DATE: 01-DEC-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 25-SEP-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 10-ACH-1992
PRIOR APPLICATION NUMBER: 07/951,747
PILING DATE: 10-ACH-1992
APPLICATION NUMBER: 07/927,337
PILING DATE: 10-AUG-1992
ATTORNEY/AGENT INFORMATION:
                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
TELEFAX: 617-428-7045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
                                                                                            TELEX:
; INFOMMATION FOR SEQ ID NO: 4
; SEQUENCE CHARACTERISTICS:
; LENGTH: 39 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLLGY: linear
US-08-461-097A-40
                                                                                                                                                                                                                                                                                                                              Best Local Similarity 10.89
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 14
US-08-466-446-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-466-446-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sklar, Robert
APPLICANT: Marchionni, Mark
APPLICANT: Gwynne, David
TITLE OF INVENTION: METHODS FOR TREATING MUSCLE
TITLE OF INVENTION: DISEASES AND DISORDERS
NUMBER OF SEQUENCES: 383
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
CITY: Boston
Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 CGCCKCCRTTCACRCAGAAGGTCTTCTCCTTCTCAGG 39
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100.0%; Score 69; DB 8;
Best Local Similarity 10.8%; Pred. No. 3.4e+03;
Matches 4; Conservative 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: Diskette
COMPUTER: DISKETTEN
OPERATING SYSTEM: DOS
SOFTWARE: FRAESEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,097A
FILING DATE: 05-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA: 08/209,204
FILING DATE: 08-MAR-1993
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                               NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04585/028001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEPRAX: 200154
                                                                 SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,097
                                                                                                                                FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: 08/209,204
FILING DATE: 08-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/059,022
FILING DATE: 06-MAY-93
ATTORNEY AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-461-097A-40; Sequence 40, Application US/08461097A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
                                           OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: MA
COUNTRY: US
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-461-097-40
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Best Local Similarity 10.8%; Pred. No. 3.40+03;
Matches 4; Conservative 33; Mismatches 0; Indels
Query Match 100.0%; Score 69; DB 8; Length 39; Best Local Similarity 10.8%; Pred. No. 3.4e+03; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: 225 FLUCTORY STATE: Boston STATE: Massachusetts
ZIP: 0211-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
MADIUM TYPE: Diskette, 5.25 inch, 360 kb storage
                                                                                                                                                                                                                       US-08-467-459-40

Sequence 40, Application US/08467459

SERREAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Gaynne, David I.; Marchionni, Mark;
APPLICANT: McBurney, Robert N.
TITLE OF INVENTION: INHIBITORS OF CELL PROLIFERATION,
TITLE OF INVENTION: THEIR PREPARATION AND USE
NUMBER OF SEQUENCES: 184

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: DISACLOE, 5.25 INCH, 550 AND COMPUTER: IBM OPERATING SYSTEM: PC-DOS SOCFWARE: WORDDERECT COURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/467,459 FILING DATE: CLASSIFICATION DATA: APPLICATION NUMBER: 08/011,396 FILING DATE: 29-JAN-1993 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/984,085 FILING DATE: 01-DEC-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/981,747 FILING DATE: 01-DEC-1992 PRIOR APPLICATION NUMBER: 07/951,747 FILING DATE: 25-SEP-1992 PRIOR APPLICATION NUMBER: 07/951,747 FILING DATE: 01-DEC-1992 ATTORNEY/AGENT INFORMATION: NAME: Clark, Paul T. REGISTRATION NUMBER: 04585/017004 TELECOMMUNICATION INFORMATION: REFERENCE/DOCKET NUMBER: 04585/017004 TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Fish & Richardson STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: 39

TYPE: amino acid
STRANDEDNESS:

TOPOLOGY: linear
US-08-467-459-40
                                                                                                                                                                                                         RESULT 15
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Search completed: May 27, 2005, 00:45:15 Job time : 496 secs

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US-10-530-253-14
US-10-530-253-28
US-10-530-253-28
US-10-530-253-28
US-60-655-875-134564
US-60-655-875-134564
US-60-655-875-134564
US-60-655-875-170437
PCT-US04-05292-65
PCT-US04-05292-65
PCT-US04-05292-65
PCT-US04-05292-65
US-10-530-253-30
US-10-530-253-30
US-10-530-253-34
US-10-703-032-106880
US-10-703-032-141430
                                                                                                                     US-11-031-175-16481
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May 27, 2005, 00:32:31; Search time 52 Seconds (without alignments) 81.197 Million cell updates/sec
                                                                                                                           537409
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Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                              537409 segs, 114114291 residues
                             - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Title: Perfect score:

Sequence:

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Scoring table:

Searched:

28, Appl 36, Appl 36, Appl 134564, 16835, 170437, 5, Appl 65, Appl

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Sequence 65, Appl Sequence 30, Appl Sequence 34, Appl Sequence 106880, Sequence 16481, A

ENE DISRUPTION, GENOME OF ACTERIUM AND GENOME CHIP USING

RESULT 1 US-10-526-324-813 ; Sequence 813, Application US/10526324	; GENERAL INFORMATION: ; APPLICANT: Imanaka, Takayuki ; APPLICANT: Imanaka, Takayuki ; TITLE OF INVENTION: METHOD OF TARGETED GENE ; TITLE OF INVENTION: HYPERTHERMOSTABLE BACTI ; TITLE OF INVENTION: THE SAME ; FILE REFERENCE: 490051.401USPC ; CURRENT APPLICATION NUMBER: US/10/526,324 ; PRIOR APPLICATION NUMBER: PCT/1B2003/003597	; PRIOR FILING DATE: 2003-08-29 ; PRIOR APPLICATION NUMBER: JP 2002-319011 ; PRIOR FILING DATE: 2002-08-30 ; NUMBER OF SEQ ID NOS: 2167 ; SOFTWARE: Patentin version 3.1 ; SEQ ID NO 813 : LENGTH: 53	TYPE: PRT ORGANISM: Thermococcus kodakaraensis KOD1	; realuks: ; NAME/KEY: misc_feature . rocartow: (414542)	OTHER INFORMATION: n is a or c or g or t.	, FEATURE: , NAME/KEY: misc_feature	; COTHER INFORMATION: n is a or c or g or t.	; FEATURE: ; NAME/KEY: misc_feature	; LOCATION: (786907)(786907) ; OTHER INFORMATION: n is a or c or q or t.		; LOCATION: (786944)(786946)	5		; UTHER INFORMATION: n is a or c or g or t. : FEATURE:	NAME/KEY: misc_feature	; LOCATION: (1128488)(1128488)	
		hance to have a sault being printed, tribution.	Description	Sequence 813, App					Sequence 26460, A Sequence 26457, A	Sequence 107111, Sequence 192031,				Sequence 164331, Sequence 123185,		Sequence 159412, Sequence 192934.	
ch 0% ch 100% st 45 summaries	nts AA New:*  todata71/paa/PCT_NEW_COMB.pep:*  todata11/paa/US06_NEW_COMB.pep:*  todata11/paa/US08_NEW_COMB.pep:*  todata11/paa/US08_NEW_COMB.pep:*  todata11/paa/US09_NEW_COMB.pep:*  todata11/paa/US09_NEW_COMB.pep:*  todata11/paa/US101_NEW_COMB.pep:*	of results predicted by chance to have a equal to the score of the result being printed, sis of the total score distribution.	סו פס	6 US-10-526-324-813 7 IIS-11-097-143-26469	, ;,	, ر, ر	. ر. د		7 US-11-097-143-26460 7 US-11-097-143-26457	6 US-10-703-032-107111 6 US-10-703-032-192031	8 US-60-669-175-32704 6 US-10-467-657-5372	, ט,	, ن ر	ر; د	υ,	6 US-10-703-032-159412 6 US-10-703-032-192934	6 US-10-703-032-105601 6 US-10-530-253-29
Minimum Match Maximum Match Listing first	Pending Patents AA  1: /cgn2_6/ptodata, 2: /cgn2_6/ptodata, 3: /cgn2_6/ptodata, 5: /cgn2_6/ptodata, 6: /cgn2_6/ptodata, 7: /cgn2_6/ptodata, 8: /cgn2_6/ptodata,	No. is the number of greater than or equal derived by analysis	Query Match Length	100.0 53	0.00		0.00			100.0 76 100.0 76					0,0	100.0	100.0 95 100.0 97
Post-processing:	Database :	Pred. No. i score great and is deri	Result No. Score	1 69 1	69		000	6 6 9 9	0 1 69	69 9	5 69	69 9	. 80	0 0 0	1 69	3 69	69

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APPLICANT: et al.

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID

TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE

TITLE OF INVENTION: DROSOPHILA GENES.

FILE REFERENCE: CL000728

CURRENT PILING DATE: 1999-10-05

PRIOR PLICATION NUMBER: 60/157,832

PRIOR PLICATION NUMBER: 60/160,191

PRIOR PLILNG DATE: 1999-10-19

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR PLILNG DATE: 1999-10-12

PRIOR PLILNG DATE: 1999-11-12

PRIOR PLILNG DATE: 1999-11-12

PRIOR PLILNG DATE: 1999-11-12

PRIOR PLILNG DATE: 1999-11-28

PRIOR PLILNG DATE: 2000-02-24

PRIOR PLILNG DATE: 2000-03-23

NUMBER: 60/191,637

PRIOR PLILNG DATE: 2000-03-23

NUMBER OF SEQ ID NOS: 43008

SSOC ID NOS: 43008

1:PNATH. FASTENCY FOR WINDOWS VERSION 4.0

1:PNATH. FASTENCY FOR WINDOWS VERSION 5.
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GENERAL INFORMATION:
Sequence 658, Application US/10526324
GENERAL INFORMATION:
Taboul APPLICANT: Imanaka, Takayuki
APPLICANT: Atomi, Haruyuki
TITLE OF INVENTION: HTPERTHERMOSTABLE BACTERIUM AND GENOME OF
TITLE OF INVENTION: HTPERTHERMOSTABLE BACTERIUM AND GENOME CHIP USING
TITLE OF INVENTION: HTPE SAME
FILE REFERENCE: 490051.401095C
CURRENT APPLICATION NUMBER: US/10/526,324
PRIOR APPLICATION NUMBER: PCT/1B2003/003597
PRIOR APPLICATION NUMBER: DCT/1B2003/003597
PRIOR APPLICATION NUMBER: JP 2002-319011
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                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 CGPCCGPCCGPCGGCGCGPCYGPNVCGPCYACGPC 46
                                                                                                                                                                                                                            3 CGPCGSCCGYYCCGPCCGPCGPRCGPCGSCCGPCGPC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 69; DB 7;
10.8%; Pred. No. 1.3e+02;
iive 33; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                               ; Sequence 26490, Application US/11097143; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Venter, J. Craig APPLICANT: et al.
                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: DROSOPHILA
; TYPE: PRT
; ORGANISM: DROSOPHILA
US-11-097-143-26469
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Best Local Similarity
Matches 4; Conserv
                                                                                                 Query Match
Best Local Similarity
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APPLICANT: Venter, J. Craig
TITLE OF INVENTION: DREADY, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
FILE REFERENCE: CLOOO728
CURRENT APPLICATION NUMBER: 60/157,832
PRIOR APPLICATION NUMBER: 60/150,932
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR PLIING DATE: 1999-10-19
PRIOR PLIING DATE: 1999-10-19
PRIOR PLLING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR APPLICATION NUMBER: 60/164,769
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR PLLING DATE: 1999-12-28
PRIOR PLLING DATE: 1999-12-28
PRIOR PLLING DATE: 2000-01-12
PRIOR PLLING DATE: 2000-01-12
PRIOR PLLING DATE: 2000-03-23
NUMBER: FALING DATE: 2000-03-23
NUMBER: FEALERSQ for Windows Version 4.0
SEQ ID NO 26469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                           ή.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc feature
LOCATION: (1561477)..(1561477)
OTHER INFORMATION: n is a or c or g or t.
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NAME/KEY: misc feature

LOCATION: (1767941) .. (1767941)

GTER INFORMATION: n is a or c or g or t.

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                                                                                                                                                                           NAME/KEY: misc feature
LOCATION: (1128517)...(1128518)
OTHER INFORMATION: n is a or c or g or
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                                 OTHER INFORMATION: n is a or c or g or
                                                                                                                                or
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                                                                          NAME/KEY: misc feature
LOCATION: (1128505)..(1128506)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1561400)..(1561400)
OTHER INFORMATION: n is a or c
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (1561545)..(1561545)
OTHER INFORMATION: n is a or c
                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (1128539)..(1128540)
OTHER INFORMATION: n is a or c
     LOCATION: (1128499)..(1128499)
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NAME/KEY: DOMAIN
LOCATION: (10)..(56)
OTHER INFORMATION: Vertebrate metallothioneins proteins domain identified by
OTHER INFORMATION: eMATRIX, accession number BL00203, p-value=9.603e-21, raw score
OTHER INFORMATION: 13.94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: Venter, J. Craig

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID

TITLE OF INVENTION: DETECTION EXPESSION OF 10,000 OR MORE

TITLE OF INVENTION: DETECTION EXPESSION OF 10,000 OR MORE

TITLE OF INVENTION: DROSOPHILA GENES.

TITLE OF INVENTION: DROSOPHILA GENES.

CURRENT PILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: 60/150,191

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR PILING DATE: 1999-10-28

PRIOR PILING DATE: 1999-11-12

PRIOR PILING DATE: 1999-11-139

PRIOR PILING DATE: 1999-11-15

PRIOR PILING DATE: 2001-01-12

PRIOR PILING DATE: 2000-01-12

PRIOR PILING DATE: 2000-01-2

PRIOR PILING DATE: 2000-01-24

PRIOR PILING DATE: 2000-01-32
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                  RESULT 5

US-10-450-763-53958

Sequence 53958, Application US/10450763

GENERAL INPORMATION:
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
CURRENT PAPPLICATION NUMBER: US/10/450,763
CURRENT PAPPLICATION NUMBER: PCT/US01/08631

PRIOR PILING DATE: 2003-06-11
PRIOR PILING DATE: 2001-03-30
PRIOR PILING DATE: 2000-03-31

PRIOR PILING DATE: 2000-03-31

PRIOR PILING DATE: 2000-03-31

PRIOR PILING DATE: 2000-08-23

NUMBER OF SEQ ID NOS: 60736

SOFTWARE: CLUSTOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 61;
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-11-097-143-26472
; Sequence 26472, Application US/11097143
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-450-763-53958
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 53958
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CVKCYICWKFCPEPAIYIKPDGYVAIDYDYCKGCGIC 43
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TYPE: PRT
ORGANISM: Thermococcus kodakaraensis KOD1
                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
LOCATION: (414542)...(414542)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1561477)..(1561477)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1128505)..(1128506)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1128517)..(1128518)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (1561545)..(1561545)
OTHER INFORMATION: n is a or c or g or t.
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LOCATION: (839139)..(839139)
OTHER INFORMATION: n is a or c or g or
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LOCATION: (786944)..(786946)
OTHER INFORMATION: n is a or c or g or
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NAME/KEY: misc_feature
LOCATION: (786890)..(786890)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128488)..(1128488)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128539)..(1128540)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1561400)..(1561400)
OTHER INFORMATION: n is a or c or g
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LOCATION: (786907)..(786907)
OTHER INFORMATION: n is a or c or g
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LOCATION: (1128499)..(1128499)
OTHER INFORMATION: n is a or c
                       NUMBER OF SEQ ID NOS: 2167
SOFTWARE: PatentIn version 3.1
PRIOR FILING DATE: 2002-08-30
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Best Local Similarity
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                                                                        SEQ ID NO 658
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Matches
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TYPE: PRT ORGANISM: DROSOPHILA

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APPLICATION GARY L. Breton
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FR
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILE REPERENCE: PATHO0-03DIV1
CURRENT APPLICATION NUMBER: US/11/079,045
CURRENT APPLICATION NUMBER: US 60/128,705
FRIOR APPLICATION NUMBER: US 60/128,705
FRIOR APPLICATION NUMBER: US 999-04-09
FRIOR APPLICATION NUMBER: 2009-04-04
FRIOR APPLICATION NUMBER: US 09/540,209
FRIOR APPLICATION NUMBER: US 09/540,404
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TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DEOSOPHILA GENES.
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                                                                                                                    Length 71;
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                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          %; Score 69; DB 7;
; Pred. No. 1.8e+02;
33; Mismatches 0
                                                                                                                 100.0%; Score 69; DB 7; 10.8%; Pred. No. 1.8e+02;
                                                                                                                                                              33; Mismatches
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CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT FILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR PILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-19

PRIOR PLING DATE: 1999-10-18

PRIOR PLING DATE: 1999-10-18

PRIOR APPLICATION NUMBER: 60/164,769

PRIOR PLING DATE: 1999-11-12

PRIOR APPLICATION NUMBER: 60/173,383

PRIOR PLING DATE: 1999-11-12

PRIOR PLING DATE: 1999-11-12

PRIOR PLING DATE: 1999-11-12

PRIOR PLING DATE: 2000-02-24

PRIOR PLING DATE: 2000-02-23

NUMBER OF SEQ ID NOS: 43008
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12-11-09-143-26460
; Sequence 26460, Application US/11097143
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 9113, Application US/11079045; GENERAL INFORMATION:
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Best Local Similarity 10.8%;
Matches 4; Conservative 33,
                                                                                              Query Match
Best Local Similarity 10.0%
Local 4, Conservative
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                   TYPE: PRT
ORGANISM: B.fragilis
US-11-079-463-9113
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US-11-079-045-9113
                                                                                                                                                                                                                                                                                                                                                          US-11-079-045-9113
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LENGTH: 71
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR CONTROL OF INSECT INFESTATION IN PLAFILE REFERENCE: 38-21(53596)
CURRENT APPLICATION NUMBER: 0560842
PRIOR FILING DATE: 2004-04-09
PRIOR FILING DATE: 2004-04-27
PRIOR APPLICATION NUMBER: 6655532
PRIOR APPLICATION NUMBER: 66579062
PRIOR FILING DATE: 2004-06-11
PRIOR APPLICATION NUMBER: 6603421
PRIOR APPLICATION NUMBER: 6603421
PRIOR FILING DATE: 2004-06-11
PRIOR APPLICATION NUMBER: 66017261
PRIOR APPLICATION NUMBER: 66017261
PRIOR FILING DATE: 2004-06-11
PRIOR FILING DATE: 2004-06-11
PRIOR FILING DATE: 2004-10-11
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GENERAL INFORMATION:
APPLICATION:
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/11/079,463
CURRENT FILING DATE: 2005-03-14
CURRENT FILING DATE: 1999-04-09
PRIOR PILING DATE: 1999-04-09
PRIOR APPLICATION NUMBER: US 60/128,705
PRIOR APPLICATION NUMBER: US 60/128,705
PRIOR APPLICATION NUMBER: US 09/540,209
PRIOR APPLICATION NUMBER: US 09/540,209
NUMBER OF SEQ ID NOS: 10444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Coding regions on vcDNA: vcDNA=SeqID_411; Strand=-; Position=985-
US-60-669-241-25327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                            100.0%; Score 69; DB 7; Length 63; 10.8%; Pred. No. 1.5e+02; tive 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 69; DB 8; Length 66; Best Local Similarity 10.8%; Pred. No. 1.6e+02; Matches 4; Conservative 33; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10 CASCTNCTNSFNLTSCLDCKNCRYRTSCESCTGCTNC 46
                                                                                                                                                                        26 CGGCGPCCGGCGPCCGPCCGGTSSFCGCGPC 62
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                                                                                                                                                                                                                                                                                                                                                                                Baum, James A
Gilberteson, Larry A
Kovalic, David K
LaRosa, Thomas J
Lu, MacJong
Munyikwa, Tichifa R. I.
Roberts, James K
                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Monsanto Technology LLC
                        Query Match
Best Local Similarity 10.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Lygus hesperus
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US-11-097-143-26472
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LENGTH: 66
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APPLICANT:
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US-60-669-175-32704
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TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Venter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
TITLE OF INVENTION: USOSOPHILA GENES.
THEN REPERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
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                                                                                                                                               100.0%; Score 69; DB 7; Length 72; 10.8%; Pred. No. 1.8e+02; Ative 33; Mismatches 0; Indels
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Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                              17 CGPCCGPCCGPCCGPCCGPCGPCGPCGPCGPC 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGPCSPCGGPCGPCGPCGPCGSCCSPCGSCCAPCGPC 45
                                                                                                                                                                                                                                 CURRENT AFFILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: 60/157,832

PRIOR FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-28

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-28

PRIOR FILING DATE: 1999-11-28

PRIOR FILING DATE: 1999-11-28

PRIOR PRIOR DATE: 1999-11-28

PRIOR PRIOR DATE: 2000-01-12

PRIOR APPLICATION NUMBER: 60/175,693

PRIOR APPLICATION NUMBER: 60/191,637

PRIOR PRIING DATE: 2000-02-24

PRIOR PRIOR DATE: 2000-03-23

NUMBER OF SEQ ID NOS: 43008

SOFTWARE: FASSEED FOR WINDOWS VERSION 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33; Mismatches
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 26460
LENGTH: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 107111, Application US/10703032; GENERAL INFORMATION:
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10.8%; P
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APPLICANT: Andersen, Scott E.
APPLICANT: Byrum, Joseph R.
APPLICANT: Conner, Timothy W.
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Masucci, James D.
Zhou, Yihua
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Best Local Similarity 10.8*
Matches 4; Conservative
                                                                                                                                                                                            Conservative
                                                                                 ORGANISM: DROSOPHILA
                                                                                                                                             Query Match
Best Local Similarity
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US-10-703-032-107111
                                                                                                     US-11-097-143-26460
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LENGTH: 74
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APPLICANT: Cao, Yongwei
APPLICANT: Cao, Yongwei
APPLICANT: Masucci, James D.
APPLICANT: Masucci, James D.
APPLICANT: Abou, Yihua
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
TITLE OF INVENTION: Plants
FILLE OF INVENTION: 19140
FILLE REFERENCE: 38-21(53374)8
CURRENT APPLICATION NUMBER: US/10/703,032
PRIOR APPLICATION NUMBER: 10/020,338
PRIOR PLILNG DATE: 2001-12-12
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Best Local Similarity 10.8%; Pred. No. 2e+02;
Matches 4; Conservative 33; Mismatches
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OTHER INFORMATION: unsure at all Xaa locations
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100.0%; Score 69; DB 6.
Best Local Similarity 13.5%; Pred. No. 2e+02;
Matches 5; Conservative 32; Mismatches
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US-10-703-032-192031
FILE REPERENCE: 38-21 (53374) B
FILE REPERENCE: 38-21 (53374) B
CURRENT APPLICATION NUMBER: US/10/703,032
CURRENT FILING DATE: 2003-11-06
PRIOR APPLICATION NUMBER: 10/020,338
PRIOR PILING DATE: 2001-12-12
NUMBER OF SEQ ID NOS: 211164
SEQ ID NO 107111
LENGTH: 76
                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_TA_1529.pep
US-10-703-032-107111
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Gilbertson, Larry A
Kovalic, David K
LaRosa, Thomas J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kovalic, David K.
APPLICANT: Andersen, Scott E.
APPLICANT: Byrum, Joseph R.
APPLICANT: Conner, Timothy W.
                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Triticum aestivum
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SEQ ID NO 192031
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APPLICANT:
APPLICANT:
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AFFILACEMENT ZUITALY, TOWNOON TOWNOON TOWNOON TITLE OF INVESTITION IN PLAINING TITLE OF INVENTION COMPOSITIONS AND METHODS FOR CONTROL OF INSECT INFESTATION IN PLAINER REFERENCE: 38-21(53597)

CURRENT APPLICATION NUMBER: 05660442

PRIOR PILING DATE: 2004-04-09

PRIOR FILING DATE: 2004-04-09

PRIOR FILING DATE: 2004-04-09

PRIOR APPLICATION NUMBER: 6056532

PRIOR APPLICATION NUMBER: 6057962

PRIOR PILING DATE: 2004-04-06-11

PRIOR FILING DATE: 2004-06-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Diabrotica virgifera
; OTHER INFORMATION: Coding regions on vcDNA: vcDNA=SeqID_12401; Strand=-; Position=59
US-60-669-175-32704
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CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR FILING DATE: 2001-02-12
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 5372
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100.0%; Score 69; DB 8;
Best Local Similarity 10.8%; Pred. No. 2.1e+02;
Matches 4; Conservative 33; Mismatches 0;
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Job time : 53 secs
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US-10-467-657-5372
; Sequence 5372, Application US/10467657
; GENERAL INFORMATION:
Lu, Maolong
Munyikwa, Tichifa R. I.
Roberts, James K
                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: 60617261
PRIOR FILING DATE: 2004-10-11
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ORGANISM: Neisseria gonorrhoeae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
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APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 40774
SEQ ID NO 32704
LENGTH: 80
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Best Local Similarity
Matches 4; Conserv
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APPLICANT:
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